

Reading Polymorphemic Dutch Compounds: Toward a Multiple Route Model of Lexical Processing

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This article reports an eye-tracking experiment with 2,500 polymorphemic Dutch compounds presented in isolation for visual lexical decision while readers' eye movements were registered. The authors found evidence that both full forms of compounds (*dishwasher*) and their constituent morphemes (e.g., *dish*, *washer*, *er*) and morphological families of constituents (sets of compounds with a shared constituent) played a role in compound processing. They observed simultaneous effects of compound frequency, left constituent frequency, and family size early (i.e., before the whole compound has been scanned) and also observed effects of right constituent frequency and family size that emerged after the compound frequency effect. The temporal order of these and other observed effects goes against assumptions of many models of lexical processing. The authors propose specifications for a new multiple-route model of polymorphemic compound processing that is based on time-locked, parallel, and interactive use of all morphological cues as soon as they become even partly available to the visual uptake system.

Keywords: morphological structure, lexical processing, eye movements, compounds

Current models of morphological processing and representation in reading have explored a wide range of logically possible architectures. Sublexical models hold that complex words undergo obligatory parsing and that lexical access proceeds via their morphemes (cf. Taft, 1991; Taft & Forster, 1975, 1976). Supralelexical models, by contrast, argue that morphemes are accessed only after the compound as a whole has been recognized (e.g., Diependaele, Sandra, & Grainger, 2005; Giraudo & Grainger, 2001). Dual route models hypothesize that full-form-based processing goes hand in hand with decompositional processing. The two access routes have been usually assumed to be independent (Allen & Badecker, 2002; Baayen & Schreuder, 1999; Frauenfelder & Schreuder, 1992; Laudanna & Burani, 1995; Schreuder & Baayen, 1995), although an interactive dual route model has been proposed as well (Baayen & Schreuder, 2000). In connectionist models such as the triangle model (Seidenberg & McClelland, 1989), morphological effects are interpreted as arising due to the convergence of orthographic, phonological, and semantic codes. What all these theories have in

common is that they were developed to explain data obtained with chronometric measures for isolated reading of bimorphemic complex words. As a consequence, they tend to remain silent about the time course of information uptake in the reading of complex words.

Establishing the temporal order of activation of full forms (e.g., *dishwasher*) of complex words and of their morphological constituents (e.g., *dish* and *washer*) is critical for adjudicating between competing models of morphological processing. The present study addresses the time course of morphological processing by considering the reading of long, polymorphemic Dutch compounds. It is important to note that current models of morphological processing offer different predictions with regard to the visual recognition of such compounds. On supralelexical models, activation of the compound's full form (diagnosed by the compound frequency effect) is expected as the initial step of lexical access. After the full form of the compound is activated, one expects to observe simultaneous activation of both the left and the right constituent (diagnosed by frequency-based properties of a constituent). On strict sublexical models, the predicted order of activation is as follows: first, the left constituent of a compound; second, its right constituent; and finally (either coinciding with activation of the right constituent or following it) the full form. The sublexical model of Taft and Forster (1976) argues that activation of the compound's left constituent is sufficient to trigger the retrieval of the compound's full form. This model predicts sequential effects of the left constituent frequency and compound frequency and no effects of the right constituent. On some dual-route models of parallel processing, one expects roughly simultaneous effects of compound frequency and left constituent frequency, because it has been argued that both routes are pursued simultaneously and independently (e.g., Baayen & Schreuder, 1999). Bertram and Hyönä (2003) have also pro-

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posed a dual-route architecture with a head start for the decomposition route in case of long compounds, which predicts early effects pertaining to the compound's left constituent followed by the compound frequency effect.

Earlier eye-tracking studies have not only confirmed the joint relevance of both constituents and full-form representations for reading posited by dual-route models (Andrews, Miller, & Rayner, 2004; Hyönä, Bertram, & Pollatsek, 2004; Zwitserlood, 1994) but have also made available more precise information about the time course of morphological processing. For instance, Hyönä et al. (2004) found that for long compounds there is early activation of the left constituent (*dish*) and later activation of the right constituent (*washer*). However, two important questions about the time course of morphological processing are as yet unresolved. First, the temporal locus of compound frequency effects remains unclear. Several eye-tracking studies of compounds (cf. Andrews et al., 2004; Bertram & Hyönä, 2003; Pollatsek, Hyönä, & Bertram, 2000) have observed effects of compound frequency for the very first fixation, but these effects failed to reach significance. Event-related potential studies of reading (Hauk & Pulvermüller, 2004; Penolazzi, Hauk, & Pulvermüller, 2007; Sereno, Rayner, & Posner, 1998) have repeatedly shown early effects of whole-word frequency (< 150–200 ms), but they focused on relatively short (four to six characters) and morphologically simplex words. An early locus for the compound frequency effect in long compounds would challenge strict sublexical accounts of morphological processing, according to which whole-word frequency effects would reflect postaccess combinatorial processes instead of tapping into early visual information uptake. Second, it is unclear whether the activation of the compound's full form precedes, follows, or coincides with the activation of the compound's constituents. The present evidence is controversial. For instance, Juhasz, Starr, Inhoff, and Placke (2003) argued—on the basis of eye-tracking, lexical decision, and naming experiments—that it is the compound's head, the last constituent to be read (e.g., *washer* in *dishwasher*), that plays the decisive role in the late stages of compound recognition, whereas the effects of the initial constituent emerge early and are weak (see, however, Juhasz, 2007). A possible reason for the dominance of the right constituent is its typical semantic convergence with the meaning of the whole compound (see also Duñabeitia, Perea, & Carreiras, 2007). These results were interpreted as support for models that argue for either coactivation of the right constituent and the full form (Pollatsek et al., 2000) or activation of the right constituent following activation of the full form (Giraudo & Grainger, 2001). Their claim contrasts with chronometric studies by Taft and Forster (1976), for example, who found evidence that the left constituent guides lexical access to a compound's meaning. Taft and Forster saw these results as evidence that a compound's full form gets activated after the left constituent of the compound receives activation.

The first aim of the present study is to address the temporal order of lexical access to the full form and the morphological constituents of compounds. In other words, we explore how soon and in what order the properties of the compound's full form and the properties of the compound's left and right constituents emerge in the timeline of compound recognition. Second, we broaden the scope of constituent processing by probing whether morphological families of constituents (sets of compounds sharing a constituent; e.g., *ice pick*, *ice cube*, *ice box*) contribute to the speed of pro-

cessing over and above properties of full forms and those of constituents as isolated words. Lexical decision studies have argued that the effects of constituent families are semantic in character and hence emerge late, at the peripheral postaccess stages of the complex word processing (e.g., De Jong, Schreuder, & Baayen, 2000). In this study we tackle the temporal locus of the effects of constituent families, using eye-tracking as a technique with a better temporal resolution than the one offered by lexical decision latencies. Third, we zoom in on the issue of independence of the full-form and decompositional processing routes claimed in some dual-route parallel processing models by considering the possibility that the effects elicited by the full-form properties might be modulated by constituent properties.

Instead of investigating bimorphemic compounds, we examined compounds with three to six morphemes. With regard to type, such polymorphemic compounds are more common in Dutch than the bimorphemic compounds that are traditionally studied in the experimental literature. For instance, perusal of the CELEX lexical database (Baayen, Piepenbrock, & Gulikers, 1995) shows that 54% of the nominal compounds have more than two morphemes. An additional dimension of morphological processing that we consider as the fourth goal of our study is the role of (freestanding and bound) morphemes deeply embedded in morphological structure (e.g., *wash-* and *-er* in *dishwasher*). Are morphemes at lower levels of morphological hierarchy recognized as independent units of meaning by the human lexical processor and used in compound identification, or are they invariably treated as parts of larger structural units (e.g., *washer*)? If, as we will argue, readers maximize their use of cues available for efficient compound identification, we may expect that the deeply embedded free and bound morphemes are used in the course of processing as well.

In what follows, we report a large regression experiment with 2,500 target compounds that combined eye-tracking of isolated word reading with lexical decision as superimposed task to ensure sufficient depth of processing. We opted for this combination because it provides detailed insight into the time course of morphological processing, and it provides sufficient statistical power. In the General Discussion, we return in detail to the methodological consequences of our decision to make use of lexical decision rather than sentential reading. Here, we restrict ourselves to noting that a parallel study presenting Finnish compounds in sentential contexts (Kuperman, Bertram, & Baayen, in press) yielded a pattern of results that is highly consistent with the morphological effects reported below. Our present experiment provides evidence that current models of morphological processing are too restrictive in their architectures and that a more flexible framework in which all opportunities for recognition are maximized (Libben, 2006) is called for.

Method

Participants

Nineteen students of the Radboud University of Nijmegen (12 women and 7 men) were paid €20 (approximately \$29) for participation in the study. All were native speakers of Dutch and reported normal or corrected-to-normal vision and right-handedness.

Apparatus

Eye movements were monitored by the head-mounted video-based EyeLink II eye-tracking device (SR Research, Ltd., Missi-

ssauga, Ontario, Canada). The average gaze position error of EyeLink II is $< 0.5^\circ$, whereas its resolution is 0.01° . Recording of the eye movements was performed on the left eye only and in the pupil-only mode. The sampling rate of recording used in this study was 250 Hz. The 17-in. (43-cm) computer monitor used for the display of the stimuli had a 60-Hz refresh rate.

Stimuli

In total, 2,500 lexical items (1,250 existing words and 1,250 nonce compounds) were included as stimuli. A list of existing polymorphemic Dutch compounds (triconstituent compounds or biconstituent compounds with at least one and at most four derivational affixes) was selected from the CELEX lexical database (Baayen et al., 1995)—for instance, *werk + gev-er* [work-giver; i.e., employer]. Additionally, a list of multiply complex nonce compounds was created by blending existing words into novel combinations (i.e., combinations that are not registered in the CELEX database); for instance, *alarmijsbaan*, composed of *alarm* [alarm] and the compound word *ijsbaan* [skating ring]. At the level of immediate constituents, the resulting targets and fillers represented a mixture of noun–noun, adjective–noun and verb–noun compounds.

The average number of morphemes per stimulus was 3.2 ($SD = 0.4$). The maximum length of a stimulus was set at 12 characters. The resulting range of 8–12 characters (mean length = 11.62, $SD = 0.74$) allowed for a tight experimental control of word length, and kept collinearity of such measures as word length and frequency and left constituent length and frequency within reasonable bounds. Stimuli were displayed one at a time in a fixed-width font (Courier New 12). With a viewing distance of about 80 cm, one character space subtended approximately 0.36° of visual angle.

Procedure

Participants were instructed to read words at their own pace. They were also informed that nonce compounds were built of existing Dutch words and were asked to evaluate the whole stimulus as an existing word or a nonword by pressing the right button (“yes” response) or the left button (“no” response) of a dual-button box. Prior to the presentation of the stimuli, the eye-tracker was calibrated using a nine-point grid that extended over the entire computer screen. Prior to each stimulus, a fixation point was presented in the central position of the screen for 500 ms. After each third stimulus, a drift correction was performed using the screen-central fixation point as a mark. After 500 ms or after the calibration was corrected, a stimulus was displayed in black lowercase characters on a white background. When one of the dual buttons was pressed, the stimulus was removed from the screen and a fixation point appeared. If no response was registered after 5,000 ms, a stimulus was removed from the screen and the next trial was initiated. Participants’ responses and response times were recorded along with their eye movements.

Stimuli were displayed centralized vertically and slightly off center horizontally such that the space between the fourth and the fifth characters of a stimulus was always at the center of the screen where the fixation point was shown. This position is closest to the preferred viewing position (the most frequent position where the eyes initially land) reported in eye movement studies for Finnish, English, and French words with the lengths that we used, mostly

12 characters (e.g., Bertram & Hyönä, 2003; McDonald & Shillcock, 2004; Vergilino-Perez, Collins, & Doré-Mazars, 2004).

The presentation order of stimuli was randomized. Stimuli were presented in two separate sessions, each consisting of three blocks. The order of presentation of the blocks and the order of the words within each block were the same for each participant (see Appendix B for a discussion of randomization procedures). For each participant, sessions were run on two different dates, whereas blocks within one session were separated by a 5- to 10-min break.

After each break the eye-tracker was calibrated again. A single session lasted 70 min at most, and the total time of the experiment lasted a maximum of 130 min.

Dependent Variables

For the analysis of the lexical decision data, we considered as dependent variables the (natural) log-transformed response times (RT) as well as the accuracy of responses (Correct).

In the eye-tracking data analysis, we selected as early measures of lexical processing the first fixation duration (FirstDur), and the subgaze duration on the compound’s left constituent (SubgazeLeft; the summed duration of all fixations on the left constituent before exiting it). As measures that tap into later stages of compound recognition, we considered subgaze for the right immediate constituent (SubgazeRight; the summed duration of all fixations on the right constituent before exiting it). Gaze duration (GazeDur) served as the global measure of processing difficulty. In this study, gaze duration was defined as the summed duration of all fixations on the target word that were completed before one of two events took place: Either the reader fixated away from the word, or the lexical decision was made.¹ All durational measures were natural log-transformed to reduce the influence of atypical outliers.

We considered several other eye-movement measures as well. These included single, second, and third fixation durations; initial fixation position; the amplitude of the first within-word saccade; the probability of a given fixation being the last one on the word; the probability of a given fixation being to the left of the previous fixation; and the total number of fixations on a word. The data patterns for these measures were in line with the ones we reported but did not offer substantial additional insight into our research questions.

Predictors

Morphological variables. The measures of morphological characteristics of stimuli included: whole word (compound) frequency (WordFreq); the word frequency of the left constituent as

¹ Note that SubgazeLeft and SubgazeRight are not strictly additive in the measure of gaze duration. In the situation where Fixation 1 is on the left constituent, Fixation 2 on the right one, and Fixation 3 on the left one, SubgazeLeft is equal to the duration of Fixation 1 and SubgazeRight to the duration of Fixation 2. The measure of gaze duration, however, would be equal to the sum of Fixations 1, 2, and 3, and could show an effect that differs in size from the sum of effects found for both subgazes. Also, we fitted the statistical models to the subgaze measures with the nonzero duration. There are words, however, in which all fixations fall on one constituent, and there is no subgaze duration for the other constituent. In such cases there is only one subgaze component contributing to the composite measure of gaze duration.

an isolated word (LeftFreq); and the word frequency for the right constituent as an isolated word (RightFreq). All these frequencies were lemma frequencies, i.e., summed frequencies of a compound word and of its inflectional variants (e.g., sum of frequencies of the singular form *newspaper*, the plural form *newspapers*, and the singular and plural genitive forms *newspaper's* and *newspapers'*). All frequency-based measures in this study, including the ones reported in the remainder of this section, were obtained from CELEX (Baayen et al., 1995; counts based on a corpus of 42 million word forms) and log-transformed to reduce the influence of outliers.

We also considered measures of morphological connectivity for the constituents of our compounds. We refer to the set of compounds that share the left (right) constituent with the target as the left (right) morphological family of that constituent (e.g., the left constituent family of *ice cream* includes *ice pick*, *ice cube*, and *ice box*). Words that appear as constituents in many compounds (i.e., have large morphological families) or in frequent compounds (i.e., have high family frequency) have been repeatedly shown across languages to elicit shorter lexical decision latencies, whether presented visually or auditorily (cf. De Jong et al., 2000; De Jong, Feldman, Schreuder, Pastizzo, & Baayen, 2002; Dijkstra, Moscoso del Prado Martín, Schulpen, Schreuder, & Baayen, 2005; Moscoso del Prado Martín, Bertram, Häikiö, Schreuder, & Baayen, 2004). Left constituent family size is also known to modulate gaze duration in interaction with semantic opacity of Finnish compounds (cf. Pollatsek & Hyönä, 2005).²

Morphological family size for the left constituents in our compounds strongly correlated with the frequencies of these left constituents as isolated words. We orthogonalized these collinear measures by fitting a regression model where left constituent family size was predicted by left constituent frequency. We then considered the residuals of this model (ResidLeftFamilySize) as our new left family size measure. It was highly correlated with the original measure ($r = .95, p < .0001$), but the effects of constituent frequency were now partialled out. Using the same procedure for the right constituent family size and frequency, we obtained ResidRightFamilySize, which again closely approximated right constituent family size ($r = .93, p < .0001$), and was orthogonal to RightFreq. We decorrelated family size and frequency for analytical clarity to be better able to assess the independent contributions of predictors (beta coefficients) to the model.

The presence of each subconstituent morpheme and its position in the morphological structure were coded by the multilevel factor Affix with the following levels: initial (for compounds with prefixed left constituents), medial (for compounds with a suffixed left constituent, an interfix, a prefixed right constituent, or any combination of these affixes), final (for compounds with suffixed right constituents), multiple (for compounds with multiple affixes³) and triconstituent (for pure triconstituent compounds with three word stems and no affixes; for the sake of analytical clarity, we excluded from our analyses 112 compounds with three word stems and further affixes). The resulting counts of stimuli representing each type of morphological complexity are summarized in Table 1.

We also considered affix productivity (AffixProd; the type count of derived words in which the affix occurs). The total number of morphemes in the compounds was included as an index of the compound's morphological complexity (Complexity).

Table 1
Counts of Compounds Partitioned by Type of Morphological Complexity

Type of complexity	Number of stimuli
Triconstituent	580
Initial	158
Medial	541
Multiple	407
Final	702

Other variables. We also considered word length (WordLength; in the range of 8–12 characters), as well as left constituent length (LeftLength). The longitudinal effect of the experimental task on the participants' behavior (e.g., fatigue or habituation as the participant works through the experiment) was estimated by means of the position of the stimulus in the experimental list (TrialNum). We also took into account the influence that carried over from trial $N - 1$ to trial N (see Baayen, Davidson, & Bates, in press; De Vaan, Schreuder, & Baayen, 2007) by considering the log-transformed response time from the trials immediately preceding the current one (RT1). Other control predictors that reached significance in codetermining either the lexical decision latencies or reading times as revealed in eye movements are presented in Appendix A. Table A1 in Appendix A lists the distributions of the continuous variables used in this study, including their ranges, and mean and median values.

Statistical Considerations

In this study we made use of mixed-effects multiple regression models with random intercepts for subject and word (and occasionally by-participant random slopes and contrasts for item-bound predictors) and the predictors introduced above as fixed effect

² For both the left and the right constituents, the alternative measure of family frequency (the summed token frequency of the members in the morphological family) consistently elicited weaker effects than family size of the respective constituents in all statistical models, in contrast to the findings of De Jong et al. (2002) for Dutch compounds. The difference in effect sizes was revealed in smaller regression (beta) coefficients for family frequencies when constituent family frequencies and family sizes were included, separately, as predictors in our statistical models. For instance, in the model for gaze duration, the regression coefficient was $-.026$ for left constituent family frequency and $-.036$ for left constituent family size. As the distinction between family size and family frequency effects is not crucial for our research questions, we do not discuss this measure further. We rather note that the entropy measure proposed by Moscoso del Prado Martín et al. (2004) may be a possible resolution for the relative impacts of the family-based alternatives.

³ We classified compounds with more than one affix at the immediate constituent boundary, such as *rov-er-s-hol* [robbers' den] as medial rather than as multiple. In other words, the category medial comprises compounds with at least one medial affix, whereas the category multiple comprises compounds with affixes at more than one position in the compound. We opted not to differentiate between compounds with different numbers of medial affixes, because the effects of these affixes considered separately were very similar across our analyses.

factors and covariates (cf. Baayen, 2008; Bates & Sarkar, 2005; Pinheiro & Bates, 2000).

Unless noted otherwise, only those fixed effects are presented below that reached significance at the 5% level in a backward stepwise model selection procedure. All random effects included in our models significantly improved the explanatory value of those models, as indicated by significantly higher values of the maximum likelihood estimate of the model with a given random effect compared to the model without that random effect (all *ps* < 0.0001 using likelihood ratio tests); for detailed treatment of random effects in mixed-effects models, see Pinheiro and Bates (2000). Below we report which predictors required random slopes in addition to the random intercepts for subject and word; see Table A7 in Appendix A.

All models were fitted and atypical outliers were identified (i.e., points that fell outside the range of -2.5 to 2.5 *SD* of the residual error). Such outliers were removed from the respective datasets (and were not used in the composite eye-movement measures), and the models were refitted to avoid distortion of the model estimates due to atypical extreme observations. Below we report statistics of those refitted models.

Due to the large number of models fitted in this study, we report in Appendix A only the full specifications of the model for lexical decision latencies for existing words and the four models for the eye movement measures (first fixation duration, subgazes for the left and the right constituent, and gaze duration).

Results and Discussion

Lexical Decision

The initial lexical decision data pool consisted of 2,500 words \times 19 participants = 47,500 trials. From this dataset we excluded one word that was misspelled, as well as the trials in which the (log) RT value fell beyond 3 *SD* from the mean. Because no participant exceeded the threshold of a 30% error rate in either nonce compounds or the existing words, none were excluded. The resulting dataset consisted of 47,206 trials, of which 41,245 were correct replies. The error rate reached 23% for existing words and 3% for

nonce compounds. Thus, in the lexical decision task participants exhibited a clear bias toward “no” responses; this does not come as a surprise, given that many of the existing compounds are fairly low-frequency words and also semantically opaque words whose meaning is conceptually difficult to construct from the individual constituents, just as is the case with many nonce compounds. For correct replies, the average lexical decision latency was 763 ms (*SD* = 246) for existing words and 801 ms (*SD* = 261) for nonce compounds.

Below, we only discuss the analysis of the lexical decision latencies for the 18,217 trials with existing compounds that were correctly identified in the lexical decision task.

Morphological variables. Column RT in Table 2 summarizes the effects of compound frequency and frequency-based measures of a compound’s constituents on the lexical decision latencies (see Table A2 in Appendix A for the full specification of the model). The column provides effect sizes for morphological predictors (see Appendix A for the explanation as to how these were computed) and *p* values for main effects, as well as indicating interactions between predictors of interest. For clarity of exposition, we leave out of the table the effects of morphemes deeply embedded in the compound structure; these are discussed separately.

Both compound frequency (WordFreq) and morpheme-based frequencies (LeftFreq, Right-Freq) and morphological connectivity measures (ResidLeftFamilySize, ResidRightFamilySize) entered into negative correlations with the RTs; that is, higher frequencies or larger families facilitated compound processing. Of these predictors, compound frequency showed the greatest effect (-96 ms). These facilitatory morphological effects are in accord with previous reports of visual lexical decision experiments with Dutch and English compounds (cf., e.g., Andrews, 1986; De Jong et al., 2000, 2002; Juhasz et al., 2003). It is interesting that compound frequency interacted with left constituent frequency in such a way that the effect of compound frequency was strongest in compounds with the low-frequency left constituents and was weaker in compounds where left constituents were relatively frequent (see Figure 1).

Table 2
Summary of Morphological Effects on Durational Measures

Predictor	RT	FirstDur	SubgazeLeft	SubgazeRight	GazeDur
LeftFreq	-32 ms (<.001)	-35 ms (<.001)	-48 ms (<.001)	<i>ns</i>	-34 ms (<.001)
Interaction with:	WordFreq (.006), Figure 1	WordFreq (.006)	WordFreq (.001)		WordFreq (.01)
ResidLeftFamSize	-43 ms (<.001)	-47 ms (<.001)	-83 ms (<.001)	-24 ms (.028)	-61 ms (<.001)
Interaction with:	RightFreq (.004), Figure 2	WordLength (.008)			RightFreq (.008)
RightFreq	-37 ms (.002)	<i>ns</i>	<i>ns</i>	-27 ms (.001)	-27 ms (<.001)
Interaction with:	ResidLeftFamSize (.004), Figure 2				ResidLeftFamSize (.008)
ResidRightFamSize	-40 ms (.001)	<i>ns</i>	<i>ns</i>	-54 ms (.001)	<i>ns</i>
WordFreq	-96 ms (<.001)	-24 ms (<.001)	-42 ms (<.001)	-47 ms (<.001)	-73 ms (<.001)
Interaction with:	LeftFreq (.006), Figure 1	LeftFreq (.01)	LeftFreq (.001)		LeftFreq (.006)

Note. Numbers in columns 2–6 show sizes of statistically significant main effects. In the case where the main effect is qualified by the interaction (e.g., WordFreq by LeftFreq), we report the numerical estimate of the main effect size (WordFreq) for the median value of the interacting term (LeftFreq). The column RT provides references to figures illustrating the interactions, which are qualitatively similar for both lexical decision latencies and reading times. Numbers in parentheses provide *p* values for the main effects and interactions. RT = reaction time; FirstDur = first fixation duration; SubgazeLeft = subgaze duration on the compound’s left constituent; SubgazeRight = summed duration of all fixations on the right constituent before exiting it; GazeDur = gaze duration; LeftFreq = the word frequency of the left constituent as an isolated word; WordFreq = whole word (compound) frequency; RightFreq = word frequency for the right constituent as an isolated word; ResidLeftFamSize = residuals of left constituent family size; ResidRightFamSize = residuals of right constituent family size.

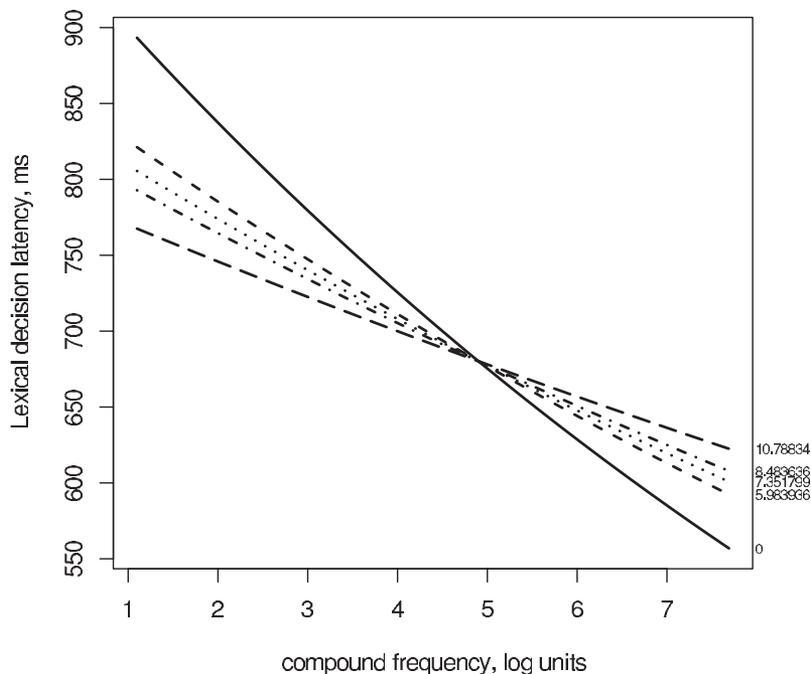


Figure 1. Interaction of compound frequency by left constituent frequency for lexical decision latencies. The lines plot the effect of compound frequency for the quartiles of left constituent frequency (quartile values provided at the right margin). Compound frequency comes with the strongest negative effect at the minimal value of left constituent frequency (solid line); the effect gradually levels off at the 1st quartile (dashed line), the 2nd quartile (dotted line), and the 3rd quartile (dot-dash line), and is weakest for the compounds with highest frequency left constituents (longdash line).

Suppose, following Libben (2006), that both compound frequency and left constituent frequency are among the morphological cues that the lexical processor may use to facilitate recognition of the compound. Then the observed interaction is the evidence that the magnitude of one such cue (e.g., left constituent frequency) appears to modulate the extent to which the other cue (e.g., compound frequency) contributes to the identification of the complex word.

We also observed an interaction between right constituent frequency and left constituent family size (see Figure 2). The effect of right constituent frequency was strongest in compounds with large left constituent families (i.e., with a large number of possible morphemic continuations for the left constituent; e.g., *shoelace*, *shoe cream*, *shoe shop*) and decreased with decreasing morphological family size.

Apparently, ease of access to the lexical representation of the right constituent (diagnosed by its frequency effect) speeds up compound recognition more when there is more uncertainty about which candidate to choose from a larger number of possible right constituents. In case the competition in the family is relatively weak due to a low number of choices, the right constituent may be relatively easy to predict, and additional morphological information in the form of right constituent frequency is not as useful for the lexical processor. Again, we find that the magnitude of one cue for compound recognition affects the utility and magnitude of other such cues.

The effects of lower-level, subconstituent morphemes revealed that compounds with two stems (of which at least one was a derivation)

were processed significantly faster than triconstituent compounds (by about 20 ms, averaged across levels of affix). Moreover, stimuli that comprised more morphemes, as measured by complexity, elicited longer latencies (effect size = 86 ms), as expected.

Other control variables. We observed habituation of participants to the task: The further they were into the experiment (as estimated by the trial position in the experimental list), the faster their lexical decisions were (effect size = -34 ms).

Longer RTs to the immediately preceding trial (RT1) went hand in hand with longer lexical decision latency at the current trial (effect size = 223 ms). These findings make a clear case that both the longitudinal effects of the experimental task and those related to immediately preceding trials contribute substantially to modulating lexical decision latencies.

Eye Movements

We considered only the first-pass reading (i.e., the sequence of fixations made before the fixation is made outside the word boundaries) and only those fixations that were completed before a response button was pressed. Trials with blinks and misreadings (i.e., trials for which no fixations were recorded by the eye-tracking device, due to machine error) were removed, as well as the trials with lexical decision latencies exceeding 3 *SD* from the mean. The resulting dataset consisted of 85,908 fixations. We also removed from the dataset of fixations and from composite eye-movement measures those fixations that exceeded 2.5 *SD* from the mean log-transformed duration, whereas the mean duration and the

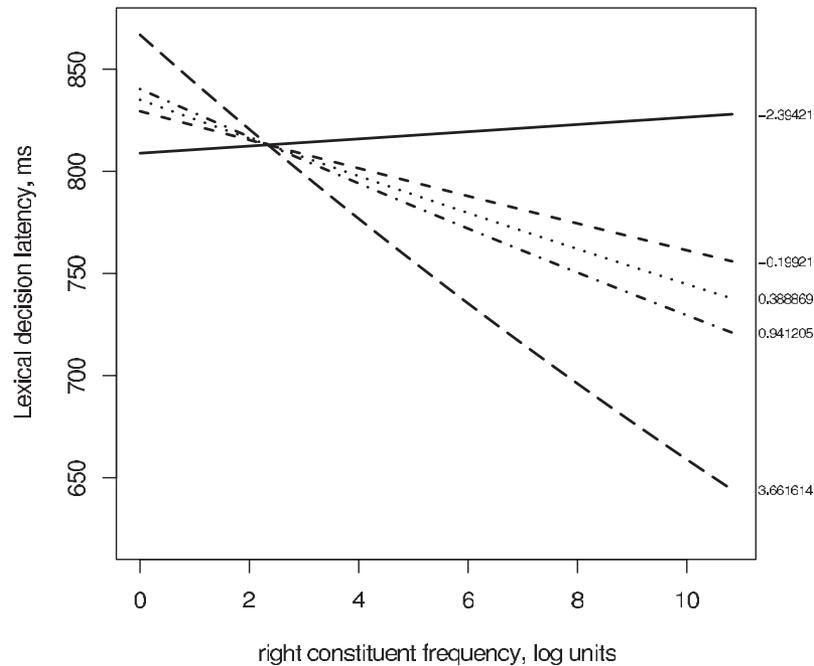


Figure 2. Interaction of right constituent frequency by (residualized) left constituent family size for lexical decision latencies. The lines plot the effect of right constituent frequency for the quartiles of left constituent family size (quartile values provided at the right margin). Right constituent frequency has no substantial effect for smallest left constituent families (solid line). The effect gradually increases at the 1st quartile (dashed line), the 2nd quartile (dotted line), and the 3rd quartile (dot-dash line), and it is strongest for compounds with the largest left constituent families (longdash line).

standard deviation were calculated separately for each participant. In this way we avoided penalizing very slow or very fast readers. In total, 2,227 (2.6%) outliers were removed, and the resulting range of fixation durations was 49 ms to 1,197 ms. Subsequently, fixations that bordered microsaccades (fixations falling within same character) were removed ($122 \times 2 = 244$ fixations, 0.1%). The resulting pool of data points consisted of 83,437 valid fixations.

Eighteen percent of the stimuli required a single fixation for reading, 36% required exactly two fixations, 26% required exactly three fixations, and it took four or more fixations to read the remaining 20% of the stimuli. The average number of fixations on a stimulus was 2.6 ($SD = 1.2$). Regressive fixations (within-word fixations located to the left of the previous fixation) constituted 12.6% of our data pool. The average fixation duration was 262 ms ($SD = 117$), and the average gaze duration was 620 ms ($SD = 382$). Eighty-one percent of initial fixations were located either on the fourth or the fifth character of the presented stimulus, which is the area where we intended those fixations to be.⁴ Seventy-seven percent of initial fixations were located on the left constituent. Because we had compounds with two- to four-character-long left constituents, a relatively large proportion of initial fixations were located at the right constituent (23%). Seventy-eight percent of second progressive fixations landed on the right constituent.

We further report our findings for the trials with existing compounds and only those that elicited correct responses. Our findings are based on four statistical models: for first fixation duration (14,232 data points), for subgaze duration on the left constituent (11,684 data

points), for subgaze duration on the right constituent (8,495 data points), and for gaze duration (14,616 data points).

Morphological effects: Compound and immediate constituents. Columns 3 to 6 in Table 2 are a summary of the effects that morphological structure elicits in eye movements across four statistical models (see full specifications for the models in Tables A3–A6 in Appendix A). Considered jointly, the results of the statistical models in Table 2 outline the temporal flow of compound recognition. First, we found evidence that both immediate constituents and the whole compound affect lexical processing of compound words (cf., e.g., Andrews et al., 2004; Bertram & Hyönä, 2003; Hyönä et al., 2004). In fact, every single morphological predictor that we considered (compound frequency, constituent frequencies, and family sizes, as well as properties of deeply embedded morphemes discussed below) had a role to play in the time course of visual compound recognition. This hints at the possibility that morpho-

⁴ It should be noted that the positions of almost 90% of initial fixations were within the measurement error ($<0.5^\circ$ of the visual angle) of EyeLink II—that is, no more than 1.4 characters away from the displayed fixation point. The shape of the distribution of initial fixation positions was close to normal with the mean of 40.7 pixels (that is, between the fourth and fifth letter) and standard deviation of 8.4 pixels. The initial fixations at the tails of the distribution (in the beginning or the end of the word) may be explained by the somewhat long presentation of the fixation point (500 ms), which may have caused people to occasionally saccade away from that fixation point prior to word presentation.

logical structure offers more cues for the task of compound identification than previously thought.

Second, properties of the left constituents of compounds showed earlier effects than the respective properties of the right constituents: the latter were only present in the late measures, Subgaze-Right and GazeDur. Moreover, the impact of the right constituent on compound recognition was considerably weaker than that of the left constituent: The effects of the right constituent were smaller in size and often qualified by interactions with other predictors. These findings may reflect the fact that the left constituent is available earlier to the lexical processor than the right constituent. The typical sequence of fixations in our dataset supported this claim: Initial fixations tended to be located at the left constituent (77% of first fixations), whereas subsequent fixations mostly landed on the right constituent (78% of progressive second fixations).⁵ We note that the size of the left constituent family code-terminated the speed of identification of a compound's right constituent. Apparently, the relative ease of processing of the left constituent spills over to the processing of the right constituent, which is consistent with the spillover effect of word N on word $N + 1$ observed in sentential reading (e.g., Rayner & Duffy, 1986; Reichle, Rayner, & Pollatsek, 2003).

Third, the compound frequency effect emerged as early as the first fixation and lingered on throughout the entire time course of compound processing. That the strong and statistically significant effect of compound frequency shows so early resolves the question raised by Bertram and Hyönä (2003, p. 627) of whether compound frequency might affect the early stages of visual processing in long compounds. The answer is that it does for 8- to 12-character-long words.⁶ The likelihood that our stimuli, which are mostly 12 characters long, were appreciated in one fixation is quite low; in fact, only 18% of our stimuli elicited a single fixation. We conclude that we have found evidence that full-form access (diagnosed by the compound frequency effect) is initiated before all characters of the compound have been foveally inspected (for the discussion of the early locus of word frequency effect, see also Cleland, Gaskell, Quinlan, & Tamminen, 2006).

Fourth, the fact that the effect of compound frequency was simultaneous with the left constituent frequency and family size effect and preceded the right constituent frequency and family size effect poses a problem for strictly sequential sublexical models of morphological processing. In such models, one would expect full-form activation to occur in time after activation of the left and the right constituent. In the Taft and Forster (1976) variant of this model, properties of the right constituent should never exert any influence on compound word identification.

Our set of findings is also problematic for supralexical models, as those models argue for initial activation of the full form and subsequent spreading activation of constituent morphemes. On this view, the properties of the left and the right constituents are expected to receive activation from the full form, and left and right constituent frequency effects should therefore kick in later than the full-form frequency effect. In fact, however, our data show that at least right constituent effects only emerge in later or global processing measures—that is, subgaze duration for the right constituent and gaze duration.

Fifth, we observed two surprising effects of constituent morphological paradigms. Left constituent family size effect showed up at the first fixation, which is unexpectedly early given the

traditional interpretation of family size effects as a postaccess semantic effect reflecting the spread of activation through morphological paradigms (cf., e.g., Bertram, Schreuder, & Baayen, 2000; De Jong et al., 2000, 2002). To explain the finding one has to assume either that the family size effect is formal rather than semantic in nature or that semantic effects can emerge earlier than is usually claimed. As we outline in the General Discussion, we believe that both the formal and the semantic components contribute to the family size effect. On the other hand, we found a late effect of ResidRightFamSize on subgaze duration for the right constituent. Recall that the right constituent family is a set of compounds (e.g., *vanilla cream*, *ice cream*, *shoe cream*, etc.) beginning in morphemes that can combine with the given right constituent (*cream*). The effect is surprising, because by the time the right constituent is scanned, it is quite plausible that the one left constituent that actually occurs in the compound (e.g., *vanilla*) has already been (partly) identified, and then activation of a paradigm of possible left constituents (e.g., *vanilla*, *ice*, *shoe*, etc.) appears unwarranted. It is likely that the effect of the right constituent family may be driven by cases in which lexical processing of the left constituent is not complete at the first fixation (e.g., due to difficult lexical processing of the left constituent or suboptimal visual uptake of word-initial information) and continues as a spillover effect even as the eyes move to the right constituent. We return to the role of morphological families in the General Discussion.

Sixth, the interactions between morphological predictors that we saw in lexical decision latencies were replicated in eye-movement measures. As early as the first fixation, left constituent frequency modulated the compound frequency effect, such that compound frequency contributed most to recognition of those compounds in which left constituent frequency was lower and the compound frequency effect diminished as the left constituent frequency increased (see Figure 1). It is important that compound frequency still has a large role to play even when the left constituent frequency is high and the traditional decompositional route is supposed to be the preferred route of compound processing. This interaction indicates that activation of compounds' full forms and of morphemes is not independent, as claimed in several dual-route models of morphological processing, and that the lexical processor

⁵ Given the lengths of our compounds and the initial fixation positions, it is likely that some characters from the right constituent are identified during an initial fixation on the left constituent. However, the absence of early effects associated with the compound's right constituent implies that the available orthographic information on the right constituent is apparently not sufficient for early activation of that morpheme (cf. Hyönä et al., 2004).

⁶ The effect of compound frequency was still significant in the statistical model for the first fixation duration from which single-fixation cases were excluded (model not shown, $p < .0001$). We did not observe an interaction of word length by compound frequency, but as the range of word lengths in our study is small, with most words having a length of 12 characters, our data do not shed light on the visual acuity hypothesis of Bertram and Hyönä (2003), according to which compound frequency effects would be more prominent for shorter words with less than 9 characters (Bertram & Hyönä, 2003; cf. also Pollatsek et al., 2000; Niswander-Klement & Pollatsek, 2006).

is not identifying compounds by strictly selecting between decomposition or full-form processing. Instead, the processing appears to be flexible and cooperative, taking advantage of both (or more, see below) routes, even when it is prompted to rely more upon one of the routes. Thus, identification of the compound through its full form is optimal when the other route is less beneficial for identification purposes and vice versa; morphological decomposition preferentially takes place when full-form access is less favorable for compound recognition. Moreover, balanced use of the two routes is in place from the earliest stages of complex word recognition.

Also, in subgaze duration for the right constituent we observed the interaction of *ResidLeftFamSize* by *RightFreq*, which showed the strongest effect of right constituent frequency in compounds with large left constituent families and thus with many potential right constituents that might follow the left constituent (see Figure 2). As we argued above, we take this interaction as evidence that morphological or other properties of morphemes and complex words serve as cues to recognition of morphologically complex structures and that some cues modulate the presence and magnitude of the effect of other cues.

Morphological effects: Deeply embedded morphemes. Thus far we have considered morphological structure at the level of the whole compound and its immediate constituents. We now consider the effects of the internal structure of these immediate constituents.

Similar to the lexical decision latencies, triconstituent compounds (i.e., those combining three lexemes) consistently elicited longer reading times in the eye-movement record than compounds with two lexemes (one of which additionally included derivational morphemes). The divergence in the processing of the two compound types did not emerge immediately at the first fixation; rather, it presented itself in subgaze and gaze durations. As effects related to meaning are assumed to occur late, we conclude that the divergence reflects a relative difficulty of semantic integration of three, rather than two, free-standing lexemes (on the temporal order of morphological and semantic effects in compounds, see, e.g., Cunnings & Clahsen, 2007).

The role of affix position in a complex word varied in accordance with the temporal order of the visual uptake. Obviously, compound-final affixes are viewed with more acuity when the compound's right constituent, rather than the left one, is under foveal inspection. Indeed, compound-final affixes elicited shorter subgaze durations and gaze durations, but their effect was five times stronger in the model for *SubgazeRight* ($\beta = -.10$, $p = .0001$) than it was in the model for *SubgazeLeft* ($\beta = -.02$, $p = .0001$). Furthermore, multiple affixes appeared to facilitate processing even more than other types of affixation, as revealed in subgaze duration for the left constituent (see Table A4). This finding is consistent with the hypothesis that affixes function as segmentation cues in locating the boundaries of morphological constituents (Kuperman et al., in press). The observed advantage of compounds with multiple affixes may indicate the relative ease of identifying a higher level morphological hierarchy in complex words with multiple segmentation cues.

An analysis of the subset of words with exactly one affix (9,790 fixations) showed that more productive affixes (i.e., affixes that occur in more word types) came with shorter gaze durations, $\beta = -.009$, $t(9790) = -6.403$, $p < .001$; effect size = -15 ms; model

not shown. This result converges with lexical decision studies in Finnish (cf. Bertram, Laine, & Karvinen, 1999) reporting shorter RTs for derived words with more productive affixes than for words with unproductive affixes.

Orthographic and visuomotor variables. Compound length (*WordLength*) went hand in hand with shorter first fixations (-37 ms) and with longer gaze durations (26 ms). This trade-off between the number and duration of fixations in correlation with word length is well attested in the eye-movement literature (cf. Vergilino-Perez et al., 2004, and references therein). Compounds with longer left constituents (*LeftLength*) elicited longer first fixations and subgaze durations for left constituents, which is as expected. In subgaze durations for the right constituents and gaze durations, the effect of left constituent length appeared to be reversed: *LeftLength* correlated negatively with durations. However, because we set the maximum for compound length, longer left constituents implied shorter right constituents. So the longer the compound's left constituent, the shorter its right constituent, and the faster it completes the visual uptake of the right constituent (hence shorter subgaze duration for the right constituent), which is in line with the direction of the corresponding effect for the left constituent length.

At first fixation, the nonlinear effect of fixation position on fixation duration showed the inverse-U shape (see the linear term *FixPos* and the quadratic term *FixPos2* in Table A3). The fixations between the 4th and the 5th character (i.e., the position of the displayed fixation point in our experiment) had a longer duration (on average by about 70 ms) than did fixations at the word's extremes, the 1st and the 12th character of the stimuli. This inverted-optimal viewing position effect is well attested in the literature on eye movements for single word recognition and sentential reading (for an overview of available theoretical accounts, see Vitu, Lancelin, & Marrier d'Unienville, 2007). Initial fixation position did not interact with any predictors of our interest.

Other control variables. We observed longitudinal effects of the course of the experiment on participants' performance. The more the participants progressed into the experiment (as measured by the position of trial in the experimental list), the shorter their first fixations were (effect size = -9 ms); their gaze durations were also shorter (effect size = -8 ms). In other words, the eye-movement record, just like the lexical decision latencies, shows that participants become familiarized with the task as the experiment proceeds, in line with, for example, Meeuwissen, Roelofs, and Levelt (2003) and De Vaan et al. (2007).

The longer the lexical decision latency to the immediately preceding trial was (*RT1*), the longer the first fixations were (effect size = 51 ms). Longer *RT1* also came with a substantial lengthening of gaze duration (effect size = 282 ms). The spillover effect on the current trial of the processing difficulty of the preceding trial is not only noticeable in the visual lexical decision latencies but also apparently codetermines the entire time course of morphological processing, starting from the first fixation onward. There may be two components to the effect of the *RT* on the preceding trial. First, this effect may reflect the spillover of the lexical processing load, which is clearly increased in the cases with longer *RT1*. In other words, word $N - 1$ may still be processed even when the lexical decision has been made and word N has been presented. Second, and perhaps more likely, the dynamics of going through the experiment may be such that the local process-

ing speed at word *N* adapts to the speed developed at previous trials (in our case, the immediately preceding trial). Being fast in a recent decision-making and motoric action of the lexical decision may influence the availability of resources and expected speed of processing for the current trial (regardless of the actual lexical characteristics of the currently presented word). We leave disentangling these possibilities to further research. Yet we note that neglecting this predictor in the statistical analysis may have profound consequences. For instance, when RT1 was removed from the statistical model for gaze durations, the amount of variance explained by the fixed effects dropped by 1.3%. From a methodological perspective, bringing longitudinal and local effects in the course of the experiment may be crucial for coming to a proper understanding of the data (cf. De Vaan et al., 2007; Kinoshita & Mozer, 2006; Taylor & Lupker, 2006).

General Discussion

This study primarily addressed the role of morphological structure in compound recognition. This section begins with a summary of findings; then, we elaborate on the methodology of this study; and finally, we formulate requirements for a model of compound processing that would account for the present set of results.

To explore computation for multiply complex words, we considered a range of diagnostic measures traditionally interpreted as indicating decompositional processing. In our data, we observed facilitatory effects of the left and right constituent lemma frequencies, as well as the facilitatory effects of the left and right constituent family sizes. In addition, we found facilitatory effects of the compound lemma frequency, the traditional hallmark for nondecompositional processing.

The time course of all these effects was tied to the time course and direction of reading. Properties associated with the left constituent played a role in the early measures of eye movements, whereas the role of the right constituent emerged relatively late (cf. Hyönä et al., 2004). Moreover, the effect sizes observed for the right constituent were considerably smaller.

The constituent frequency and family size effects may have arisen at the level of form processing, at the level of semantic processing, or possibly at both levels. At the level of form, the effect of a constituent's frequency may reflect the reader's experience with identifying that constituent's string of characters. The effect of morphological family may tap into a reader's more specific experience with parsing out and recognizing the constituent as part of a larger word. At the level of word meaning, a constituent's frequency may gauge the ease of access to its meaning. A constituent's family size would then estimate the resonance that activation of a constituent morpheme gives rise to in its morphological family.

The effect of compound frequency emerged already at the first fixation duration, a point when most compounds have not yet been fully scanned. There are several ways in which this surprising effect can be interpreted. This full-form frequency effect may result from unstructured form processing in which the available visual input at the first fixation (the initial characters, the previewed characters in the middle of the word, as well as the word's length; cf. Pollatsek & Rayner, 1982; Rayner, Well, Pollatsek, & Bertera, 1982) is matched against stored form representations. The more entrenched this full-form representation is, the earlier the

benefits of its availability emerge in the eye-movement record. It is important to note that this interpretation presupposes that full-form representations do not require full visual inspection of the input and may be accessed on the basis of partially matching information (cf. de Almeida & Libben, 2002). The fact that the effect of compound frequency is also visible in later measures implies that the full-form representation of a compound is actively involved in the process of compound recognition even when other sources of lexical information become available, possibly for checking the new input for consistency with the already activated full form and/or deactivating other competitors in the morphological family.

It is unlikely, however, that unstructured form processing would fully account for the compound frequency effect and especially for its presence in the late eye-movement measures. The compound frequency effect survives inclusion in the statistical model of the frequency of the initial quadrogram summed over words that match the target compound in length (model not shown). This indicates that it is unlikely that the compound frequency effect can be reduced specifically to the earliest available visual information. Following Baayen, Wurm, and Aycocock (2007), it is conceivable that full-form frequency effects reflect, at least in part, memory traces of constituent morphemes having been combined together into one lexical unit. The higher the frequency of a complex word in language, the stronger the association between that word and its morphemes, and the more experience the reader has with integrating a given morpheme into that embedding word. If so, a high-frequency compound may benefit more from identification of one of its constituents than a low-frequency compound.

At the present stage of our knowledge, we cannot exclude that the compound frequency effect is also indicative of facilitation from semantic processing, given that semantic effects have been observed for very short initial time spans (cf., e.g., Diependaele et al., 2005; Hauk & Pulvermüller, 2004; Penolazzi et al., 2007; cf. also Baayen, Feldman, & Schreuder, 2006, for evidence concerning a strong semantic component to the word frequency effect).

In addition to constituent frequency and family size effects and the compound frequency effect, we obtained ample evidence for a role of morphemes that are embedded inside the immediate constituents of compounds. Thus, embedded affixes that are more productive elicited shorter gaze durations, as expected given previous studies of bimorphemic derivations (e.g., Bertram et al., 1999). We also observed that compounds embedded in compounds require more reading time than derivations embedded in compounds. We have two possible explanations for that. First, compounds with three free-standing lexemes are more difficult to integrate semantically than those with two such lexemes. For instance, readers need to determine whether a compound with three lexemes is left-branching (i.e., the first two constituents modify the third, as in *voet-bal + bond* [football association]) or right-branching (i.e., the first constituent is a modifier of the two latter constituents, as in *zaal + voet-bal* [indoor football]). Second, the derivational morpheme may have served as a parsing cue to identification of immediate constituents, and using such cues allows faster access to morphological constituents and faster semantic wrap-up of the complex word (see Kuperman et al., in press, for a more detailed discussion of this issue).

Methodological Considerations

A comparison of the results obtained with the visual lexical decision task and those obtained with the cumulative eye-movement measures (subgaze and gaze durations) show remarkable convergence. In the RTs, just as in eye movements, we observe facilitatory effects of constituent frequencies and family sizes and also those of compound frequencies. We also find qualitatively similar interactions between morphological predictors (WordFreq by LeftFreq, and ResidLeftFamSize by RightFreq) in lexical decision latencies and eye-movement durational measures. Furthermore, embedded morphemes and experimental control variables give rise to very similar patterns of results in the two datasets, lexical decision latencies and eye movements. What the analysis of the eye movements adds is detailed information about the time course of morphological processing, including the early and lingering compound frequency effect, the early left constituent family size effect, and the temporal sequence of the effects pertaining to the compounds' left and right constituents.

Our choice of investigating the processing of isolated existing and nonce compounds in visual lexical decision has offered us both advantages and disadvantages. The main advantage of using isolated words is the ability to collect large numbers of data points from the same participant relatively quickly. As a result, our statistical analyses enjoy the benefit of enhanced power. In addition, combining lexical decision, the task that has been used most intensively to study morphological processing, with eye tracking allows us to evaluate to what extent the two paradigms converge (cf. the program of investigating functional overlap between tasks described in Grainger, 2003). As noted above, there is indeed remarkable convergence in our data.

Our choice for using isolated words in lexical decision also comes with several disadvantages, most of which concern the issue of the ecological validity of our results. In single word reading, there is no parafoveal preview from the preceding word, and there is no natural spillover effect from the target word to the next word to be investigated. More important, lexical decision may induce rather different kinds of processing strategies than those used for the natural integration of word meaning into the sentence and discourse.

Another methodological decision that we had to make is whether to include a look-away point on the screen—that is, whether to instruct participants to complete their lexical decision task by fixating on either the word *Yes* or the word *No*, which would be displayed in two different areas on the screen equally distant from the area where the target word was displayed (for the full description of this technique, see Hyönä, Laine, & Niemi, 1995). For compatibility with the existing body of literature, we stayed as close to the conventional lexical decision paradigm as possible and did not make use of such a look-away point. Instead, we considered in our analyses only those fixations that were completed before the button press registering a lexical decision. The price we pay is the possibility of some more noise in the eye-movement measures, especially in the gaze durations. Yet in our data, gaze durations and RTs are not that highly correlated: $R^2 = .46$ only. Thus, both gaze durations and RTs serve as dependent variables in their own right.

We also note that the presence of nonce compounds and many low-frequency existing compounds in our experiment may have enhanced decompositional processing and inhibited full-form pro-

cessing. In the light of this possibility, it is all the more surprising that an effect of compound frequency is observed at the very first fixation.

Whatever the disadvantages of our methodology may be, the pattern of results that we have obtained and reported either in the body of the article or in Appendix A dovetails perfectly with many of the results obtained in the literature for sentential reading, such as visuo-oculomotor effects (cf., e.g., O'Regan, Lévy-Schoen, Pynte, & Brugailière, 1984; Rayner, 1998; Vitu, McConkie, Kerr, & O'Regan, 2001); effects of compound length and frequency, as well as of constituent frequencies (cf., e.g., Andrews et al., 2004; Duñabeitia et al., 2007; Hyönä et al., 2004; Hyönä & Pollatsek, 1998; Juhasz et al., 2003; Taft & Forster, 1976); and effects of orthographic *n*-grams (reported in Appendix A; cf. Lima & Inhoff, 1985). Furthermore, in a recent sentential reading study (Kuperman et al., in press) in which Finnish compounds were embedded in context, a highly similar pattern of results was observed, including early effects of compound frequency, left constituent frequency, and family size; later and weaker effects of right constituent frequency and family size; and interactions between morphological predictors, as well as longitudinal experimental effects.

Toward a Theory of Compound Processing

According to Libben (2006), readers and listeners maximize their opportunities for comprehension by the simultaneous use of all processing cues available to them and all processing mechanisms that they have at their disposal, including retrieval from memory and compositional computation. The present study provides support for Libben's hypothesis of maximization of opportunity. All constituent morphemes, the whole compound itself, and morphological families that share one of the compound's constituents play a noticeable role in lexical processing of compounds. This indicates that there are multiple routes at work in compound processing and readers use these routes interactively, at different times and to a different extent, to efficiently and accurately recognize compounds. The early compound frequency effect shows that readers do not wait to see all the characters of the word before making inferences about the word's identity. The early compound frequency effect also shows that readers do not gain access to compound representations only after having accessed the compounds' constituents. The interactions of morphological predictors (compound frequency by left constituent frequency and left constituent family size by right constituent frequency) show that the cues modulate each other and that decompositional processes and full-form-driven processes are not independent. Using one kind of morphological information for compound identification as if other sources of information do not exist amounts to missing out on the cumulative use of information and on concomitant facilitation of performance.

In what follows, we take as the point of departure the basic assumption of parallel dual-route models, given the evidence in our data for both processing routes. As the detailing of a full-fledged model of morphological processing is beyond the scope of this study, we restrict ourselves to listing a number of requirements that are not satisfied by the current parallel dual-route models proposed in the literature (e.g., Schreuder & Baayen, 1995). Although our results were obtained in the visual domain, we believe

that the requirements outlined below would equally hold for the models of the auditory processing of compounds.

First, current models of morphological processing almost always discuss complex words as if they are read with only one fixation. An example of a model that addresses the temporal dynamics of reading complex words is the one proposed by Polatsek, Reichle, and Rayner (2003); they conclude that a parallel dual-route architecture is unable to approximate the empirical data unless the two routes of lexical processing are allowed to interact. It is clear, also from the present data, that the details of the time course of information entering the system need to be explicitly included in models of morphological processing in reading.

In the typical left-to-right reading of long compounds, the very first opportunities for comprehension of the compound already present themselves during parafoveal preview, when information about the initial characters and word length becomes available (Rayner et al., 1982). In single-word reading, this information is also available very early, during the low-level attentional scan of the word that occurs in the beginning of fixation (cf. Reichle et al., 2003). Following Clark and O'Regan (1999) and O'Regan (1979), word length may play a disambiguating role in word recognition (for the opposing view, see Inhoff & Eiter, 2003). For words embedded in the sentential context, additional information may come from contextual predictability (e.g., Ehrlich & Rayner, 1981), collocational strength (e.g., McDonald & Shillcock, 2001) and constructional cues (e.g., Frazier, Carminati, Cook, Majewski, & Rayner, 2006).

The next opportunities for restricting the range of possible interpretations for the visual input arise at the first fixation, where a range of properties of the first constituent come into play: not only the frequency of the left constituent, its length, and its morphological family, but also the combinatorial likelihood of morphemes within the whole compound in conjunction with information about the compound's length. Later opportunities (at second and subsequent fixations) include properties of the right constituent. New information obtained at this stage is processed against the backdrop of the information already extracted about the word.

Second, models of morphological processing in reading need to allow for a simultaneous processing of information at different levels without requiring strict sequentiality of processing stages; witness, for instance, the simultaneous early effects in our data of compound frequency, left constituent frequency and family size, and orthographic *n*-gram effects.⁷ Our results challenge sublexical models, which allow full-form access only after morphological constituents have been recognized (cf. Pinker, 1999; Taft, 1991, 2004; Taft & Forster, 1976). Our results also challenge supralexical models, which only allow constituents to come into play after the compound as a whole has been recognized (Giraudo & Grainger, 2001).

Third, models of compound processing should allow for the modulation of the weight of one opportunity by the presence and strength of other opportunities, as demonstrated by the interaction of compound frequency and left constituent frequency (for a related discussion of cue trade-offs in speech processing, see, e.g., Mattys, White, & Melhorn, 2005; McClelland & Elman, 1986). Current parallel dual-route models tend to simplify morphological processing to activation of autonomous lexical representations that are blind to each other's activation (cf. Frauenfelder & Schreuder,

1992; Laudanna & Burani, 1985; Schreuder & Baayen, 1995; see, however, Baayen & Schreuder, 2000). In general, the fact that we also find in the parallel study early constituent frequency effects and whole-word frequency effects at the same time tells us that one cue or route is not canceling out the other completely, a prediction that would directly derive from a strict dual-route model. Depending on the strength of the available cues, the fine-tuning of this kind of cooperative system depends on the specific properties of the complex word.

Fourth, models of morphological processing should come to grips with fast activation of morphological paradigms (families) associated with a compound's constituents. One important constraint on morphological models is our finding that left and right constituent families are activated immediately upon access to those constituents and not after full-form access.

Effectively, a model that meets these requirements is no longer a dual-route model but rather a multiple-route model that, in morphological terms, allows access to full forms, immediate constituents, embedded morphemes, and morphological families. More generally, such a model will have as its basic principle maximization of all opportunities—morphological, orthographic, phonological, and contextual—for comprehension of the visual input. We believe that probabilistic and information-theoretical approaches to lexical processing developed recently in morphological and syntactic research (cf., e.g., Levy, 2008; Moscoso del Prado Martín et al., 2004) hold promise for formalization of those opportunities and for computational implementation of the multiple-route model of compound recognition.

⁷ A modeling framework that may prove to be useful here is the hierarchical temporal memory framework proposed by Hawkins and George (2006; see also Hawkins & Blakeslee, 2004). In the hierarchical temporal memory framework, the simultaneous processing would be accomplished by generation skip—that is, lower-level detectors in the hierarchy propagating information about the input to higher levels, skipping intermediate levels.

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(Appendixes follow)

Appendix A

Key to Table A1

Predictors of primary interest for this study are presented in the main body of the article. Additional control variables that show significant effects in our statistical models are as follows: *Correct1*, the binary indicator of whether the previous trial was a correct lexical decision; *FixPos* and *FixPos2*, first fixation position and its squared value; *FinTrigram*, frequency of the word-final trigram; and *Nomore*, indicator of whether the fixation is word-final. In addition to these, we considered a large number of control variables that were not significant predictors of reading times, fixation probabilities, or lexical decision latencies. These included variables listed in *Dependent Variables* as well as initial trigram frequency, mean bigram frequency of the word, position of the minimal bigram, affix length, branching of triconstituents, and frequencies of deeply embedded stems in triconstituents.

Specifications of Statistical Models (Tables A2–A7)

Specifications include estimates of the regression coefficients; 95% highest posterior density intervals (HPDs), which are a Bayesian estimate of the most likely values of a parameter, roughly comparable to traditional 95% confidence intervals; *p* values estimated by the Monte Carlo Markov chain (MCMC) method, using

1,000 simulations; and *p* values obtained with the *t* test for fixed effects using the difference between the number of observations and the number of fixed effects as the upper bound for the degrees of freedom (see Pinheiro & Bates, 2000, for discussion of the method). We also report the estimated standard deviations for each random intercept (e.g., Subject or Word) and each random slope (e.g., Subject by WordLength), together with the estimates based on the MCMC samples and HPD intervals, such as the MCMC mean and 95% HPDs (see Table A7 for all models); see Pinheiro and Bates (2000) for detailed treatment of random effects in mixed-effects models.

Computation of Effect Sizes (Tables A2–A6)

Effect sizes were estimated as follows. For factors, for which we used contrast coding, effect size was calculated as the difference between (a) the sum of the intercept and the contrast coefficient, β , and (b) the intercept. For log-transformed dependent variables (fixation duration, gaze duration, RT), effect sizes were calculated for back-transformed values, so that effect sizes are reported in milliseconds. Effect sizes for simple main effects of a covariate were calculated as the difference between the model's predictions for the minimum and maximum values of that covariate.

Table A1
Summary of Continuous Variables

Variable	Range (Adjusted range)	<i>M</i> (<i>SD</i>)	<i>Mdn</i>
RT	270–2,208 ms (5.6–7.7 log units)	6.7 (0.3)	6.6
InitPos	0.1–11.9 characters (1–119 pixels)	40.7 (8.4)	40.0
FirstDur	50–1,200 ms (3.9–7.1 log units)	5.6 (0.4)	5.6
SubgazeLeft	60–1,808 ms (4.1–7.5 log units)	5.8 (0.5)	5.7
SubgazeRight	82–1,097 ms (4.4–7.0 log units)	5.6 (0.4)	5.5
GazeDur	50–2,208 ms (3.9–8.2 log units)	6.5 (0.5)	6.5
TrialNum	1–2,500	12.0 (7.2)	12.0
RT1	148–4,023 ms (5.0–8.3 log units)	6.73 (0.3)	6.7
WordLength	8–12 characters	11.6 (0.7)	12.0
LeftLength	2–10 characters	5.4 (1.6)	5.0
FinTrigram	1–984,609 (0–13.8 log units)	9.6 (2.6)	9.9
WordFreq	3–2,207 (1.1–7.7 log units)	2.2 (1.1)	1.9
LeftFreq	1–24,343 (0.0–10.1 log units)	5.0 (2.9)	5.4
RightFreq	1–49,020 (0–10.8 log units)	4.5 (3.0)	4.2
ResidLeftFamilySize	3–298 (–2.3–3.7)	0.0 (1.0)	0.0
ResidRightFamilySize	3–270 (–3.5–7.4)	0.0 (1.1)	–0.1
AffixProd	3–6,002 (0.7–8.7 log units)	6.8 (1.3)	6.99
Complexity	3–6 morphemes	3.2 (0.4)	3.0

Note. Numbers in the second column show original value ranges for predictors. If any transformations have been made with the original values for statistical reasons (i.e., natural log transformation, decorrelation with other predictors, or scaling), the numbers in the parentheses show the ranges actually used in statistical models. Means, standard deviations, and median values refer to the predictor values used in the models. Values for frequency and family size measures are based on the corpus with 42 million word forms. RT = reaction time; InitPos = initial fixation position; FirstDur = first fixation duration; SubgazeLeft = summed duration of all fixations on the left constituent before exiting it; SubgazeRight = summed duration of all fixations on the right constituent before exiting it; GazeDur = gaze duration; TrialNum = position of the stimulus in the experimental list; LeftLength = length of left constituent; FinTrigram = frequency of the word-final trigram; WordFreq = whole word (compound) frequency; LeftFreq = the word frequency of the left constituent as an isolated word; RightFreq = word frequency for the right constituent as an isolated word; ResidLeftFamSize = residuals of left constituent family size; ResidRightFamSize = residuals of right constituent family size; AffixProd = affix productivity.

Table A2
Fixed Effects of the Model for Lexical Decision RT for Existing Compounds

Variable	Estimate	MCMC <i>M</i>	HPD95lower	HPD95upper	<i>p</i> MCMC	<i>pr</i> (> <i>t</i>)
Intercept	5.9740	5.9771	5.8176	6.1336	.001	.0000
WordLength	0.0148	0.0149	0.0083	0.0226	.002	.0000
LeftFreq	-0.0181	-0.0183	-0.0250	-0.0115	.001	.0000
RightFreq	-0.0095	-0.0096	-0.0129	-0.0060	.001	.0000
Complexity	0.0639	0.0634	0.0302	0.0953	.001	.0002
Trial	-0.0041	-0.0042	-0.0050	-0.0032	.001	.0000
RT1	0.1288	0.1286	0.1144	0.1413	.001	.0000
Correct1Y	-0.0160	-0.0159	-0.0285	-0.0031	.012	.0146
ResidLeftFamSize	0.0114	0.0111	-0.0106	0.0353	.354	.3557
ResidRightFamSize	-0.0122	-0.0121	-0.0194	-0.0049	.001	.0010
AffixFinal	-0.0527	-0.0526	-0.0796	-0.0295	.001	.0001
AffixInitial	-0.0178	-0.0169	-0.0613	0.0339	.500	.4801
AffixMedial	-0.0382	-0.0378	-0.0653	-0.0116	.006	.0062
AffixMultAffix	-0.0897	-0.0887	-0.1371	-0.0394	.001	.0004
WordFreq	-0.0717	-0.0722	-0.0904	-0.0533	.001	.0000
LeftFreq:WordFreq	0.0037	0.0037	0.0012	0.0062	.002	.0047
RightFreq:ResidLeftFamSize	-0.0049	-0.0048	-0.0079	-0.0017	.001	.0040

Note. RT = reaction time; MCMC = Monte Carlo Markov chain; HPD95lower = lower boundary of the 95% highest posterior density interval; HPD95upper = upper boundary of the 95% highest posterior density interval; *p*MCMC = *p* values estimated by the MCMC method using 1,000 simulations; *pr*(> |*t*) = *p* values obtained with the *t* test using the difference between the number of observations and the number of fixed effects as the upper bound for the degrees of freedom; LeftFreq = the word frequency of the left constituent as an isolated word; RightFreq = word frequency for the right constituent as an isolated word; RT1 = reaction time in the previous trial; Correct1Y = level Y of the binary indicator of whether the previous trial was a correct lexical decision; ResidLeftFamSize = residuals of left constituent family size; ResidRightFamSize = residuals of right constituent family size; AffixFinal, AffixInitial, AffixMedial, AffixMultAffix = levels of the factor Affix, which codes the positions of subconstituent morphemes; WordFreq = whole word (compound) frequency.

Table A3
Model for First Fixation Duration

Variable	Estimate	MCMC <i>M</i>	HPD95lower	HPD95upper	<i>p</i> MCMC	<i>pr</i> (> <i>t</i>)
Intercept	5.8489	5.8532	5.5993	6.1337	.001	.0000
NomoreTRUE	0.2345	0.2356	0.1773	0.3067	.001	.0000
WordLength	-0.0394	-0.0390	-0.0575	-0.0202	.001	.0000
LeftLength	-0.0261	-0.0260	-0.0304	-0.0209	.001	.0000
FixPos	0.0088	0.0088	0.0065	0.0113	.001	.0000
FixPos2	-0.0001	-0.0001	-0.0001	0.0000	.001	.0000
WordFreq	-0.0347	-0.0346	-0.0490	-0.0171	.001	.0001
LeftFreq	-0.0172	-0.0172	-0.0231	-0.0115	.001	.0000
ResidLeftFamSize	0.0728	0.0733	0.0001	0.1559	.058	.0690
Trial	0.0000	0.0000	0.0000	0.0000	.001	.0000
RT1	0.0352	0.0348	0.0168	0.0515	.001	.0001
WordFreq:LeftFreq	0.0031	0.0031	0.0010	0.0052	.010	.0058
WordLength:ResidLeftFamSize	-0.0085	-0.0086	-0.0156	-0.0019	.018	.0171

Note. MCMC = Monte Carlo Markov chain; HPD95lower = lower boundary of the 95% highest posterior density interval; HPD95upper = upper boundary of the 95% highest posterior density interval; *p*MCMC = *p* values estimated by the MCMC chain method using 1,000 simulations; *pr*(> |*t*) = *p* values obtained with the *t* test using the difference between the number of observations and the number of fixed effects as the upper bound for the degrees of freedom; NomoreTRUE = level TRUE of the factor Nomore; LeftLength = length of left constituent; FixPos = fixation position; WordFreq = whole word (compound) frequency; LeftFreq = the word frequency of the left constituent as an isolated word; ResidLeftFamSize = residuals of left constituent family size; RT1 = reaction time in the previous trial.

(Appendixes continue)

Table A4
Model for Subgaze Duration for the Left Constituent

Variable	Estimate	MCMC <i>M</i>	HPD95lower	HPD95upper	<i>p</i> MCMC	<i>pr</i> (> <i>t</i>)
Intercept	5.9312	5.9380	5.7110	6.1295	.001	.0000
WordLength	-0.0628	-0.0627	-0.0729	-0.0529	.001	.0000
LeftLength	0.0777	0.0774	0.0687	0.0856	.001	.0000
WordFreq	-0.0591	-0.0598	-0.0846	-0.0341	.001	.0000
LeftFreq	-0.0272	-0.0275	-0.0361	-0.0175	.001	.0000
RightFreq	-0.0028	-0.0029	-0.0070	0.0015	.184	.2056
ResidLeftFamSize	-0.0378	-0.0382	-0.0472	-0.0280	.001	.0000
ResidRightFamSize	-0.0023	-0.0024	-0.0116	0.0062	.624	.6280
Trial	0.0000	0.0000	0.0000	0.0000	.004	.0026
RT1	0.0748	0.0747	0.0529	0.0988	.001	.0000
AffixMedial	-0.0472	-0.0468	-0.0831	-0.0151	.008	.0084
AffixFinal	-0.0216	-0.0218	-0.0588	0.0138	.266	.2556
AffixMultAffix	-0.0805	-0.0808	-0.1153	-0.0472	.001	.0000
WordFreq:LeftFreq	0.0052	0.0053	0.0017	0.0085	.001	.0019

Note. MCMC = Monte Carlo Markov chain; HPD95lower = lower boundary of the 95% highest posterior density interval; HPD95upper = upper boundary of the 95% highest posterior density interval; *p*MCMC = *p* values estimated by the MCMC method using 1,000 simulations; *pr*(> |*t*) = *p* values obtained with the *t* test using the difference between the number of observations and the number of fixed effects as the upper bound for the degrees of freedom; LeftLength = length of left constituent; WordFreq = whole word (compound) frequency; LeftFreq = the word frequency of the left constituent as an isolated word; RightFreq = word frequency for the right constituent as an isolated word; ResidLeftFamSize = residuals of left constituent family size; ResidRightFamSize = residuals of right constituent family size; AffixFinal, AffixMedial, AffixMultAffix = levels of the factor Affix, which codes the positions of subconstituent morphemes; RT1 = reaction time in the previous trial.

Table A5
Model for Subgaze Duration for the Right Constituent

Variable	Estimate	MCMC <i>M</i>	HPD95lower	HPD95upper	<i>p</i> MCMC	<i>pr</i> (> <i>t</i>)
Intercept	5.6285	5.6278	5.3264	5.9335	.001	.0000
WordLength	0.0289	0.0289	0.0139	0.0408	.001	.0000
LeftLength	-0.1105	-0.1105	-0.1211	-0.0994	.001	.0000
WordFreq	-0.0340	-0.0339	-0.0444	-0.0244	.001	.0000
LeftFreq	-0.0016	-0.0016	-0.0068	0.0037	.550	.5612
RightFreq	-0.0103	-0.0103	-0.0167	-0.0043	.001	.0009
ResidLeftFamSize	-0.0154	-0.0153	-0.0296	-0.0027	.028	.0292
ResidRightFamSize	-0.0188	-0.0188	-0.0317	-0.0074	.001	.0052
Trial	0.0000	0.0000	0.0000	0.0000	.040	.0528
RT1	0.1029	0.1032	0.0676	0.1364	.001	.0000
AffixMedial	-0.0598	-0.0594	-0.1149	-0.0120	.020	.0210
AffixFinal	-0.1022	-0.1016	-0.1513	-0.0529	.001	.0000
AffixMultAffix	-0.0005	-0.0010	-0.0520	0.0486	.966	.9852

Note. MCMC = Monte Carlo Markov chain; HPD95lower = lower boundary of the 95% highest posterior density interval; HPD95upper = upper boundary of the 95% highest posterior density interval; *p*MCMC = *p* values estimated by the MCMC method using 1,000 simulations; *pr*(> |*t*) = *p* values obtained with the *t* test using the difference between the number of observations and the number of fixed effects as the upper bound for the degrees of freedom; LeftLength = length of left constituent; WordFreq = whole word (compound) frequency; LeftFreq = the word frequency of the left constituent as an isolated word; RightFreq = word frequency for the right constituent as an isolated word; ResidLeftFamSize = residuals of left constituent family size; ResidRightFamSize = residuals of right constituent family size; AffixFinal, AffixMedial, AffixMultAffix = levels of the factor Affix, which codes the positions of subconstituent morphemes; RT1 = reaction time in the previous trial.

Table A6
Gaze Duration Model

Variable	Estimate	MCMC <i>M</i>	HPD95lower	HPD95upper	<i>p</i> MCMC	<i>pr</i> (> <i>t</i>)
Intercept	5.6415	5.6385	5.4218	5.8684	.001	.0000
WordLength	0.0173	0.0173	0.0032	0.0319	.012	.0174
LeftLength	-0.0173	-0.0173	-0.0291	-0.0061	.002	.0029
WordFreq	-0.0912	-0.0908	-0.1194	-0.0621	.001	.0000
LeftFreq	-0.0253	-0.0253	-0.0354	-0.0142	.001	.0000
RightFreq	-0.0080	-0.0081	-0.0132	-0.0026	.008	.0054
ResidLeftFamSize	0.0073	0.0075	-0.0291	0.0446	.672	.6975
ResidRightFamSize	-0.0070	-0.0072	-0.0177	0.0025	.164	.2102
Trial	0.0000	0.0000	-0.0001	0.0000	.001	.0000
RT1	0.1506	0.1509	0.1273	0.1752	.001	.0000
AffixMedial	-0.0812	-0.0798	-0.1201	-0.0400	.001	.0001
AffixFinal	-0.1001	-0.0985	-0.1413	-0.0585	.001	.0000
AffixMultAffix	-0.0834	-0.0826	-0.1250	-0.0470	.001	.0001
FinTrigram	-0.0070	-0.0071	-0.0124	-0.0015	.012	.0185
RightFreq: ResidLeftFamSize	-0.0068	-0.0068	-0.0119	-0.0018	.008	.0087
WordFreq:LeftFreq	0.0053	0.0053	0.0016	0.0091	.008	.0055

Note. MCMC = Monte Carlo Markov chain; HPD95lower = lower boundary of the 95% highest posterior density interval; HPD95upper = upper boundary of the 95% highest posterior density interval; *p*MCMC = *p* values estimated by the MCMC method using 1,000 simulations; *pr*(> |*t*) = *p* values obtained with the *t* test using the difference between the number of observations and the number of fixed effects as the upper bound for the degrees of freedom; LeftLength = length of left constituent; WordFreq = whole word (compound) frequency; LeftFreq = the word frequency of the left constituent as an isolated word; RightFreq = word frequency for the right constituent as an isolated word; ResidLeftFamSize = residuals of left constituent family size; ResidRightFamSize = residuals of right constituent family size; AffixFinal, AffixMedial, AffixMultAffix = levels of the factor Affix, which codes the positions of subconstituent morphemes; FinTrigram = frequency of the word-final trigram; RT1 = reaction time in the previous trial.

Table A7
Random Effects for RT, FirstDur, SubgazeLeft, SubgazeRight, and GazeDur

Estimate	<i>SD</i>	MCMC <i>M</i>	HPD95lower	HPD95upper
Lexical decision latency				
Word	0.095	0.095	0.090	0.101
Subject	0.151	0.155	0.110	0.215
Residual	0.241			
First fixation duration				
Word	0.049	0.050	0.042	0.057
Subject	0.42	0.415	0.286	0.608
Subject by Nomore	0.099	0.098	0.070	0.148
Subject by WordLength	0.035	0.035	0.024	0.051
Residual	0.289			
Subgaze duration for the left constituent				
Word	0.088	0.087	0.078	0.096
Subject	0.114	0.12	0.087	0.167
Residual	0.335			
Subgaze duration for the right constituent				
Word	0.010	0.097	0.075	0.116
Subject	0.107	0.110	0.079	0.158
Residual	0.456			
Gaze duration				
Word	0.014	0.122	0.113	0.132
Subject by LeftLength	0.015	0.014	0.008	0.023

(Appendixes continue)

Table A7 (continued)

Estimate	SD	MCMC M	HPD95lower	HPD95upper
Subject by WordLength	0.017	0.018	0.012	0.025
Subject	0.082	0.022	0.001	0.172
Residual	0.386			

Note. RT = reaction time; FirstDur = first fixation duration; SubgazeLeft = summed duration of all fixations on the left constituent before exiting it; SubgazeRight = summed duration of all fixations on the right constituent before exiting it; GazeDur = gaze duration.

Comparison between effect sizes of numeric variables obtained in our study and those obtained in previous studies that set these variables to a discrete number of levels for factorial designs is not straightforward. Our estimates are defined over the entire range of values of the variable, whereas a factorial contrast is defined as a difference between group means, where groups are formed (in the

simplest case) by dichotomization of a given predictor. The best approximation to factorial estimates is one half of our effect sizes, which is equivalent (for linear effects) to the factorial contrast where the variable of interest is dichotomized and where the group means are positioned at the first and the third quartiles. Obviously, factors do not pose such a problem and are directly comparable across reports.

Appendix B

In the present study, we chose to present readers with a fixed order of items in the block and the fixed order of blocks, so that each reader saw the words in the same order (even though that one order was set randomly). We hypothesized that by using one list order we would have tighter experimental control, especially as we have the position of an item in the experimental list as a covariate in the model, so that longitudinal effects of practice or fatigue are modeled explicitly. By using the fixed list order we also attempted to avoid the increase in between-subjects variance, which derives from the random ordering of items across participants. By that, we aimed at gaining increased statistical power. Using the fixed list order, however, goes against the common practice of counterbalancing (or otherwise randomizing) the presentation order of items across participants. The problem that is usually claimed to follow from using the fixed item order is that the variance that one attributes to a predictor of interest might in fact be due to the influence of the item order. In other words, the item order is a potential confound for estimates of other effects.

It turns out that the item order is no a priori reason for worry. Linear mixed effects models are very well able to disentangle the various sources of variance for a design such as we used. In a simulation study, we considered a repeated measures design with 20 participants, 1,000 items, list position (the rank or trial number in the list) as a predictor, and five predictors specifying properties linked to the items (standing for word length, word frequency, left constituent family size, and left constituent and right constituent frequencies). In other words, the simulated data have the same design as our experimental data, albeit with fewer predictors and fewer items. The question of interest is whether the mixed-effects modeling algorithm can adequately separate the different sources of variance under two conditions, one in which each participant is exposed to the items in the same order (as in our article) and one in which each participant is exposed to the items in a different random order. This simulation does not aim to assess the significance of our predictors or to validate our statistical models. Rather, we simulate two types of experimental designs (with a fixed number of items, participants, and predictors) to

see whether, under these conditions, predictions of statistical models would differ across designs.

More formally, let i index participants, j index items, and k index trial number. Furthermore, let X_1 denote trial, X_2 to X_6 item-bound properties, and β_{0-6} the intercept and regression coefficients, respectively. We further denote participant random effect as b_{Si} (normally distributed with standard deviation σ_S) and item random effect as b_{Wj} (normally distributed with standard deviation σ_W), and we denote the error term as ϵ_{ijk} (normally distributed with standard deviation σ). Our simulated data have the general form

$$y_{ijk} = \beta_0 + \beta_1 X_{1k} + \beta_2 X_{2j} + \beta_3 X_{3j} + \beta_4 X_{4j} + \beta_5 X_{5j} + \beta_6 X_{6j} + b_{Si} + b_{Wj} + \epsilon_{ijk}, \quad b_{Si} \sim N(0, \sigma_{Si}), \\ b_{Wi} \sim N(0, \sigma_{Wi}), \quad \epsilon_{ijk} \sim N(0, \sigma). \quad (1)$$

In building a simulation there are many choices to be made. In this simulation we make a simplifying assumption that item-bound predictors X_2 to X_6 are uncorrelated, although the predictors in the empirical data show mild collinearity. Also, it is not necessarily the case that there is a unique value for a predictor for each trial. Say that if we take word length to range from 4 to 12 characters, there will not be 1,000 different values of word length for 1,000 trials; rather, integer values for 4 to 12 will be repeated multiple times, just as in the original data.

We distinguish between a model with fixed order for all participants, so that $k = j$, henceforth $M_{k=j}$, and a model in which each participant has a different order, so that $k \neq j$, henceforth $M_{k \neq j}$. We studied the behavior of both models for 20 participants and 1,000 items across 1,000 simulation runs. Columns 1–3 in Table B1 specify which fixed and random predictors were used in the simulation, what their coding is in Equation 1, and what the values were that we set for those predictors. We based the ranges of values for item-bound predictors on the actual ranges in the experimental data. Values of X_2 to X_6 varied randomly (uniformly) in the corresponding value ranges. For all predictors, only integer values were considered. Our estimates for regression coefficients and the intercept closely follow the output of the statistical model

for lexical decision latencies (Table A2), with a few exceptions. We increased variance in data by setting higher values for random errors, and we diminished the influence of the strongest lexical predictors, word length and word frequency, by dividing their regression coefficients by 10. We increased noise and weakened some effects because the simulation run on the original data showed almost perfect accuracy in estimating the coefficients, and it reported significance of predictors correctly in almost 100% of cases. Results of the simulation are summarized in Table B1.

Columns 4 and 6 in Table B1 show the means of the estimates of the coefficients for the fixed effects and for the standard deviations of the random effects, obtained with the model with fixed order of items and the model with the randomized order of items for each participant, correspondingly. Columns 5 and 7 show proportions of correctly reported significance across simulation runs for both types of models. It is evident that for large data samples, such as we used in this study, there is no appreciable difference across presentation orders in the performance of statistical models, neither in the accuracy of estimates for model coefficients or for standard deviations of random effects, nor in the power to detect the effect of the item-bound predictors. For smaller samples, we have seen cases where a single experimental list (fixed order) comes with slightly reduced power than the list with the random presentation of items.

We have carried out more simulations with different values for the fixed and random effect parameters; time and again the pattern is like the one summarized in Table B1. It is important to note that these simulations show no a priori reason to believe that sources of variance are confounded—at least, not with the number of items and the number of uncorrelated predictors that we used here. It is important to realize that the strength of linear mixed-effects models lies precisely in their ability to “unconfound” different sources of variance.

We have double-checked whether there were interactions of this longitudinal effect with item-bound predictors, including lexical, distributional, or orthographic characteristics of compounds as whole words or their morphemes, but there were none. This gives additional assurance that the morphological effects of our primary interest are not modulated by the longitudinal effects of the experimental list. We have also investigated whether other longitudinal effects might be present (ranging from priming effects due to constituents that appeared earlier in the list to effects of sharing onset or rhyme). None turned out to be significant. In other words, the morphological and orthographic effects that we report are not artifacts or confounds of experimental control variables, as can be demonstrated both in a simulation and in an experiment.

Table B1
Parameters, Estimates of the Parameters, and Power ($\alpha = .05$) for the Models Without ($M_{k = j}$) and With ($M_{k \neq j}$) Random Orders of Items for Each Participant

Predictor	Parameter	Value	$M_{k = j}$		$M_{k \neq j}$	
			Estimate	Power	Estimate	Power
Intercept	β_0	5.82	5.8074	1.00	5.8084	1.00
Trial	β_1	-0.0048	-0.0010	1.00	-0.0010	1.00
WordFreq	β_2	-0.0043	-0.0045	0.35	-0.0045	0.35
WordLength	β_3	0.0013	0.0010	0.05	0.0010	0.05
LeftFamSize	β_4	-0.0084	-0.0082	0.27	-0.0082	0.27
LeftFreq	β_5	-0.0026	-0.0027	0.21	-0.0027	0.22
RightFreq	β_6	-0.0072	-0.0072	0.88	-0.0072	0.88
Subject	σ_s	0.32	0.3100		0.3100	
Item	σ_w	0.20	0.1999		0.1999	
Residual	Σ	0.60	0.5996		0.5996	

Note. Averages over 1,000 simulation runs. WordFreq = whole word (compound) frequency; LeftFamSize = left constituent family size; LeftFreq = the word frequency of the left constituent as an isolated word; RightFreq = word frequency for the right constituent as an isolated word.

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