Analytical choices for analyzing multidimensional behavior - Many analysts test hypotheses about human speech.

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- Writing Original Draft Preparation, Writing Review & Editing; Second Author: Writing
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- 13 Editing.

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Abstract

One or two sentences providing a **basic introduction** to the field, comprehensible to a scientist in any discipline.

Two to three sentences of **more detailed background**, comprehensible to scientists in related disciplines.

One sentence clearly stating the **general problem** being addressed by this particular study.

One sentence summarizing the main result (with the words "here we show" or their equivalent).

Two or three sentences explaining what the **main result** reveals in direct comparison to what was thought to be the case previously, or how the main result adds to previous knowledge.

One or two sentences to put the results into a more **general context**.

Two or three sentences to provide a **broader perspective**, readily comprehensible to a scientist in any discipline.

Keywords: crowdsourcing science, data analysis, scientific transparency, speech, acoustic analysis

Word count: X

Analytical choices for analyzing multidimensional behavior - Many analysts test hypotheses about human speech.

1 Introduction

In order to effectively accumulate knowledge, science needs to (i) produce data that 37 can be replicated using the original methods and (ii) arrive at robust conclusions 38 substantiated by the data. In recent coordinated efforts to replicate published findings, the 39 scientific disciplines have uncovered surprisingly low success rates for (i) (Camerer et al., 2018; e.g., Open Science Collaboration, 2015) leading to what is now referred to as the replication crisis. Beyond the difficulties of replicating scientific findings, a growing body of evidence suggests that the theoretical conclusions drawn from data are often variable even when researchers have access to reliable data (REFS). The latter situation has been referred to as the inference crisis (Rotello, Heit, & Dubé, 2015; Starns et al., 2019) and is, among other things, rooted in the inherent flexibility of data analysis (Gelman & Loken, 2014; often referred to as researcher degrees of freedom: Simmons, Nelson, & Simonsohn, 2011). Data analysis involves many different steps, such as inspecting, organizing, transforming, and modeling the data, to name a few. Along the way, different methodological and analytical choices need to be made, all of which may influence the final interpretation of the data. These researcher degrees of freedom are both a blessing and a curse. 51 They are a blessing because they afford us the opportunity to look at nature from 52 different angles, which, in turn, allows us to make important discoveries and generate new hypothesis (e.g., Box, 1976; De Groot, 2014; Tukey, 1977). They are a curse because idiosyncratic choices can lead to categorically different interpretations, which eventually find their way into the publication record where they are taken for granted (Simmons, Nelson, & Simonsohn, 2011). Recent projects have shown that the variability between different data analysts is vast. This variability can lead independent researchers to draw different conclusions about the same data set as demonstrated by several projects

crowd-sourcing analysis strategies (Botvinik-Nezer et al., 2020; e.g., Silberzahn et al., 2018;
Starns et al., 2019). These projects, however, might still underestimate the extent to which
analysts vary because data analysis is not merely restricted to statistical inference. Human
behavior is complex and offers many ways to be translated into numbers. This is
particularly true for fields that draw conclusions about human behavior and cognition from
multidimensional data like audio or video data. In fields working on human speech
production, for example, researchers need to make numerous decisions about what to
measure and how to measure it. This is not trivial given the temporal extension of the
acoustic signal and its complex structural composition. Not only can decisions about
measuring the signal influence downstream decisions about statistical modeling, but
statistical results or modeling issues can also lead researchers to go back and revise earlier
decisions about the measuring process itself.

In this article, we investigate the variability in analytic choices when many analyst teams analyze the same speech production data, a process that involves both decisions regarding the operationalization of a complex observed signal and decisions regarding the statistical modeling. Specifically, we report the impact of the analytic pipeline on research results obtained by XX teams who gained access to the same set of acoustic recordings in order to answer the same research question.

78 1.1 Researcher degrees of freedom

Data analysis comes with many decisions like how to measure a given phenomenon or behavior, what data to submit to statistical modeling and which to exclude in the final analysis, or what inferential decision procedure to apply. However, if these decisions during data analysis are not specified in advance, we might stumble upon seemingly meaningful patterns in the data that are merely statistical flukes. This can be problematic because humans show cognitive biases that can lead to erroneous inferences. Humans filter the world in irrational ways (e.g., Tversky & Kahneman, 1974), seeing coherent patterns in

randomness (Brugger, 2001), convincing themselves of the validity of prior expectations ("I knew it," Nickerson, 1998), and perceiving events as being plausible in hindsight ("I knew 87 it all along," Fischhoff, 1975). In connection with an academic incentive system that rewards certain discovery processes more than others (Koole & Lakens, 2012; Sterling, 1959), we often find ourselves exploring many possible analytical pipelines, but only reporting a select few. This issue is particularly amplified in fields in which the raw data 91 lend themselves to many possible ways to measure (Roettger, 2019). Combined with a wide variety of methodological and theoretical traditions as well as varying levels of statistical training across subfields, the inherent flexibility of data analysis might lead to a vast plurality of analytic approaches that can lead to different scientific conclusions. Consequently, there might be many published papers that present overconfident interpretations of their data based on idiosyncratic analytic strategies (Gelman & Loken, 2014; e.g., Simmons, Nelson, & Simonsohn, 2011). These interpretations are either associated with an unknown amount of uncertainty or lend themselves to alternative interpretation if analyzed differently. However, instead of being critically evaluated, 100 scientific results often remain unchallenged in the publication record. Despite recent efforts 101 to improve transparency and reproducibility (REFS) and freely available and accessible 102 infrastructures such as provided by the Open Science Framework (osf.io, ADD), critical 103 re-analyses of published analytic strategies are still not very common because data sharing 104 remains rare (Wicherts, Borsboom, Kats, & Molenaar, 2006). 105

While this issue has been widely discussed both from a conceptual point of view
(Nosek & Lakens, 2014; Simmons, Nelson, & Simonsohn, 2011; Wagenmakers, Wetzels,
Borsboom, Maas, & Kievit, 2012) and its application in individual scientific fields
(e.g. Wichert et al. 2015, Charles et al. 2019, Roettger, 2019), there are still many
unknowns regarding the extent of analytical plurality in practice. Recent collaborative
attempts have started to shed light on how different analysts tackle the same data set and
have revealed a large amount of variability.

13 1.2 Crowdsourcing alternative analyses

In a collaborative effort, Silberzahn et al. (2018) let twenty-nine independent analysis
teams address the same research hypothesis. Analytical approaches and consequently the
results varied widely between teams. Sixty-nine percent of the teams found support for the
hypothesis, and 31% did not. Out of the 29 analytical strategies, there were 21 unique
combinations of covariates. Importantly, the observed variability was neither predicted by
the team's preconceptions about the phenomenon under investigation nor by peer ratings
of the quality of their analyses. The authors results suggest that analytic plurality is a fact
of life and not driven by different levels of expertise or bias. Similar crowd-sourced studies
recruiting independent analyst teams showed similar results.

While these projects show a large degree of analytical flexibility with impactful consequences, they dealt with flexibility in inferential or computational modeling. In these studies the data sets were fixed and data collection or measurement could not be changed.

However, in many fields the primary raw data are complex signals that need to be 126 operationalized according to the research question. In social sciences, the raw observations 127 correspond to human behavior, sometimes measured as a complex visual or acoustic signal. 128 Decisions about how to measure a theoretical construct related to that behavior or its 129 underlying cognitive processes might interact with downstream decisions about statistical 130 modeling and vice versa (Flake & Fried, 2019). To understand how analytical flexibility 131 manifests itself in a scenario where a complex decision procedure is involved in 132 operationalizing and measuring complex signals, the present paper looks at an 133 experimentally elicited speech data set.

$_{55}$ 1.3 Operationalizing speech

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Research on speech is at the heart of the cognitive sciences, informing psychological models of language, categorization, and memory, guiding methods for diagnosis and

therapy of speech disorders, and facilitating advancement in automatic speech recognition and speech synthesis. One major challenge in the speech sciences is the mapping between 139 communicative intentions and their physical manifestation. 140

Speech is a complex signal that is characterized by structurally different acoustic landmarks distributed throughout different temporal domains. Thus, choosing how to measure a phenomenon of interest is an important and non-trivial analytical decision. Take 143 for example the following sentence in 1:

(1) "I can't bear another meeting on zoom."

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Depending on the speaker's intention, this sentence can be said in different ways. If, 146 for instance, the speaker is exhausted by all their meetings, the speaker might acoustically 147 highlight the word "another" or "meeting." If, on the other hand, the speaker is just tired 148 of video conferences, they might acoustically highlight the word "zoom." 149

If we want to compare the speech signal associated with these two intentions, how do 150 we quantify the difference between them? What do we measure and how do we measure it? 151 Given the continuous and transient nature of speech, identifying speech parameters and 152 temporal domains becomes a non-trivial task. Utterances stretch over hundreds of 153 milliseconds and contain several levels of linguistically relevant units such as phrases, 154 words, syllables, and individual sounds. The researcher is thus confronted with a 155 considerable number of parameters and combinations thereof to choose from. 156

Speech categories are inherently multidimensional and dynamic: they consist of a 157 cluster of parameters that are modulated over time. The acoustic parameters of one category are usually asynchronous, i.e. they appear at different time points in the unfolding 159 signal, and overlap with parameters of other categories [e.g., Jongman, Wayland, and 160 Wong (2000); Lisker (1986); Summerfield, 1984; Winter (2014). A classical example is the 161 distinction between voiced and voiceless stops in English (i.e. /b/ and /p/ in bear vs pear). 162

This voiced/voiceless contrast is manifested by many acoustic features which can differ 163 depending on several factors, such as position of the consonant in the word and 164 surrounding sounds (Lisker, 1977). Furthermore, correlates of the contrast can even be 165 found away from the consonant, in temporally distant speech units. For example, the 166 initial l of the English words led and let is affected by the voicing of the final consonant 167 (/t, d/) (Hawkins & Nguyen, 2004). The multiplicity of phonetic cues grows exponentially 168 if we look at larger temporal domains as is the case for suprasegmental aspects of speech. 169 For example, studies investigating acoustic correlates of word stress (e.g. the difference 170 between *insight* and *incite*) have been using a wide variety of measurements, including 171 temporal characteristics (duration of certain segments or sub-segmental intervals), spectral 172 characteristics (intensity, formants, and spectral tilt), and measurements related to 173 fundamental frequency (f0) (e.g., Gordon & Roettger, 2017).

Moving onto the expression of higher-level functions like information structure and 175 discourse pragmatics, relevant acoustic cues can be distributed throughout even larger 176 domains, such as phrases and whole utterances (e.g., Ladd, 2008). Differences in position, 177 shape, and alignment of pitch modulations over multiple locations within a sentence are 178 correlated with differences in discourse functions (e.g. Niebuhr et al., 2011). The latter can also be expressed by global vs local pitch modulations (Van Heuven, Haan, Gussenhoven, 180 & Warner, 2002), as well as acoustic information within the temporal or spectral domain 181 (e.g., Van Heuven & Van Zanten, 2005). Extra-linguistic information, like speaker's 182 intentions, levels of arousal or social identity, are also conveyed by broad-domain 183 parameters, such as voice quality, rhythm, and pitch (Foulkes & Docherty, 2006; Ogden, 184 2004; White, Payne, & Mattys, 2009). 185

When testing hypotheses on speech production data, researchers are faced with many choices and possibilities. The larger the functional domain (e.g. segments vs words vs utterances), the higher the number of conceivable operationalizations. For example, when comparing two realization of example (1) (here repeated as 2), one of which is intended to

signal emphasis on another and one of which emphasizes zoom.

(2a) "I can't bear ANOTHER meeting on zoom." (2b) "I can't bear another meeting on ZOOM."

In text ref to Figure 1

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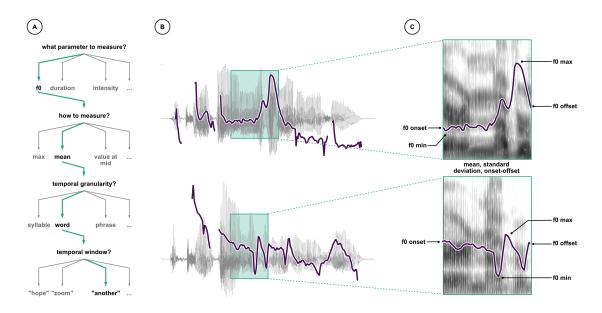


Figure 1. Illustrations of the analytical flexibility associated with acoustic analyses. (A) an example of multiple possible and justifiable decisions when comparing to utterances; (B) waveform and fundamental frequency (f0) track of two instances of utterances 2a and 2b. The words "another is highlighted by the green box"; (C) spectrogram and f0 track of the word another, exemplifying different operationalizations of differences in pitch.

Do we only compare the word another in 2a and 2b or also the word zoom or do we
measure utterance wide acoustic profiles? Do we measure the whole word? Or just the
stressed syllable? Do we average the domain or do we measure a specific point in time? Do
we measure fundamental frequency or intensity? When looking at phrase-wide temporal
domains, the number of possible analytical pipelines quickly explodes. This plurality of
analytical paths is illustrated in figure X. When comparing two utterance such as 2a and
2b, there are many things to consider. Even if we know that we want to compare

fundamental frequency of only the word another across utterances 2a and 2b, there are still many decisions to be made, all of which can be justified. For example, we could measure for 202 at specific points in time like the onset of the window, the offset, the midpoint. We could 203 also measure the value or time of the minimum or maximum f0 value. We could summarise 204 for across the entire window and extract the mean, median or standard deviation of for And 205 the garden of forking paths does not stop here. In Figure X, we went with a specific option 206 to automatically calculate fo, INSERT SOME EXAMPLES OF PITCH TRACKING 207 OPTIONS. Moreover, knowing that theses estimates are somewhat noisy, we could smooth 208 these contours to different degrees, automatically or manually remove estimates that are 200 off, etc. 210

These decisions are usually made prior to any statistical analysis, but are at times 211 revised a posteriori (i.e. after data collection and/or preliminary analyses) in light of 212 unforeseen or surprising outcomes. These myriads of possible decisions are exponentiated 213 by researcher degrees of freedom related to statistical analysis [e.g. wicherts2016]. Even the 214 analysis of a single measure can be approached via an ever-increasing range of different 215 statistical models (REFs). The present paper probes this garden of forking paths in the 216 analysis of speech. To assess the variability in data analysis pipelines across independent 217 researchers, we provided XX analytical teams with an experimentally elicited speech corpus 218 and asked them to investigate acoustic differences related to a functional contrast that 219 might be manifested across the whole utterance. 220

221 1.4 The data set - The prosody of redundant modifiers

Our data set was collected in order to answer the following research question: Do
speakers acoustically modify utterances to signal atypical word combinations? (e.g. a blue
banana vs a yellow banana)? We are interested in the acoustic profile of referring
expressions. Referring is one of the most basic and prevalent uses of language and one of
the most widely researched areas in language science. It is an open question how speakers

choose a referring expression when they want to refer to a specific entity like a banana. The context within which an entity occurs (i.e., with other non-fruits, other fruits, or other 228 bananas) plays a large part in determining the choice of referring expressions. Generally, 229 speakers aim to be as informative as possible to uniquely establish reference to the 230 intended object, but they are also resource-efficient in that they avoid redundancy (Grice, 231 1975). Thus one would expect the use of a modifier, for example, only if it is necessary for 232 disambiguation. For instance, one might use the adjective yellow to describe a banana in a 233 situation in which there is a yellow and a less ripe green banana available, but not when 234 there is only one banana to begin with. 235

Despite this coherent idea that speakers are both rational and efficient, there is much 236 evidence that speakers are often over-informative: Speakers use referring expressions that 237 are more specific than strictly necessary for the unambiguous identification of the intended 238 referent (Rubio-Fernández, 2016; Sedivy, 2003; Westerbeek, Koolen, & Maes, 2015), which 239 has been argued to facilitate object identification and making communication between 240 speakers and listeners more efficient (Arts, Maes, Noordman, & Jansen, 2011; Paraboni, 241 Van Deemter, & Masthoff, 2007; Rubio-Fernández, 2016). Recent findings suggest that the 242 utility of a referring expression depends on how good it is for a listener (compared to other referring expressions) to identify a target object. For example, Degen, Hawkins, Graf, Kreiss, and Goodman (2020) showed that modifiers that are less typical for a given referent (e.g. a blue banana) are more likely to be used in an over-informative scenario (e.g. when there is just one banana). This account, however, has mainly focused on content selection 247 (Gatt, Gompel, Deemter, & Krahmer, 2013), i.e. whether a certain referential expression is chosen or not, ignoring the fact that speech communication is much richer. 249

Even looking at morphosyntactically identical expressions, speakers can modulate
these via suprasegmental acoustic properties like temporal and spectral modifications of
the segments involved (e.g., Ladd, 2008). Most prominently, languages use intonation to
signal discourse relationships between referents (among other functions). Intonation marks

discourse-relevant referents for being new or given information to guide the listeners' interpretation of incoming messages. In many languages, speakers can use particular pitch 255 movements to signal whether a referent has already been mentioned and is therefore 256 referred back to, or a referent is newly introduced into the discourse. Many languages use 257 intonation in order to signal if a referent is contrasting with one or more alternatives that 258 are relevant to the current discourse. Content selection aside, in a scenario in which a 250 speaker wants to refer to a banana when there is also a pear on the table, the speaker 260 would most likely produce a high rising pitch accent on banana to indicate the contrastive 261 nature of the noun. In a scenario in which the speaker wants to refer to a yellow banana 262 when there is also a less ripe green banana on the table, the speaker would most likely 263 produce a high rising pitch accent on yellow to indicate the contrastive nature of the 264 modifier. In addition to a pitch accent, elements that are new and/or contrastive are often produced with additional suprasegmental prominence, i.e. segments are hyperarticulated, resulting in longer, louder and more clearly articulated acoustic targets.

268 1.5 Research questions

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The present project examines the extent to which subjective choices by different researchers analyzing a complex speech data set affect the reported results. We are further interested in which factors affect the researchers' final results.

2 Methods

We are closely following the methodology proposed by Parker et al. (Stage 1 in-principle accepted) in terms of data collection. However, the analysis will substantially diverge from their approach (see §#.#)

This project involves a series of steps (X-X):

1. We will recruit independent groups of researchers to analyze the data.

- 278 2. We will give researchers access to the speech corpus and let them analyze the data as
 they see fit.
- 3. We will ask reviewers to generate peer review ratings of the analyses based on methods (not results).
- 4. We will evaluate the variation among the different analyses.
- 5. Lastly, we will collaboratively produce the final manuscript.

We estimate that this process, from the time of an in-principle acceptance of this

Stage 1 Registered Report, will take XX months (Table X). The factor most likely to delay

our time line is the rate of completion of the original set of analyses by independent groups

of scientists.

⁸⁸⁸ 2.1 Step 1: Recruitment and Initial Survey of Analysts

The initiating authors (SC, JC, TR) created a publicly available document providing 289 a general description of the project (LINK) and a short prerecorded slide show that summarizes the study and research question in order to increase accessibility to potential 291 analysts (LINK). The project will be advertised via Social Media, using mailing lists for 292 linguistic and psychological societies (full scope of these lists is not fixed but will include 293 LIST OF LISTS), and via word of mouth. The target population is active speech science 294 researchers with a graduate degree (or currently studying for a graduate degree) in a 295 relevant discipline. Researchers can choose to work independently or in a small team. For 296 the sake of simplicity, we refer to single researcher or small teams as "analysis teams." 297

Recruitment for this project is ongoing but we aim for a minimum of XX analysis
teams independently evaluating each data set (see sample size justification below). We will
simultaneously recruit volunteers to peer-review the analyses conducted by the other
volunteers through the same channels. Our goal is to recruit a similar number of
peer-reviewers and analysts, and to ask each peer reviewer to review a minimum of four

analyses. If we are unable to recruit at least half the number of reviewers as analysis
teams, we will ask analysts to serve also as reviewers (after they have completed their
analyses). All analysts and reviewers will share co-authorship on this manuscript and will
participate in the collaborative process of producing the final manuscript. All analysts will
sign a consent (ethics) document (LINK).

To identify the minimal sample size, we followed the method in [ECO RR]. The aim of the meta-analysis is to obtain an estimate of heterogeneity of the effect sizes reported by the analysis teams (τ^2 , i.e. the variance $\sigma_{\alpha_t}^2$, see Section 2.4.2). Ideally, the 95% credible interval (CrI) of τ^2 should not include 0 (i.e. the probability p that the 95% CrI contains 0 should be less than 0.05). The probability p that a CrI interval does not include 0 is obtained via the t-statistics:

$$t = \frac{\tau^2}{SE(\tau^2)}$$

Assuming that the underlying distribution of effect sizes is normal (Knight, 2000), the standard error of τ^2 can be calculated with the formula:

$$SE(\tau^2) = \sqrt{\frac{2\tau^4}{(n-1)}}$$

where n is the sample size. Since we know p and τ^2 , we can calculate n such that p < 0.05. Plugging $SE(\tau^2)$ into the formula of the t-statistics shows that, when n is fixed, t (and hence p) will be the same regardless of τ^2 :

$$t = \frac{\tau^2}{SE(\tau^2)} = \frac{\tau^2}{\sqrt{\frac{2\tau^4}{(n-1)}}} = \sqrt{\frac{(n-1)}{2}}$$

In other words, the minimum sample size n needed to exclude 0 from the 95% CrI of τ^2 is invariant regardless of the estimate of heterogeneity τ^2 . When n=12 then $t_{(12-1)}=t_{(11)}=2.3452$ and p=0.0388, which is below the 0.05 threshold, as required. In

sum, a minimal sample of 12 effect sizes (i.e. of 12 analysis teams) would thus be sufficient to exclude 0 from the 95% CrI of τ^2 .

324 2.2 Step 2: Primary Data Analyses

The analysis teams will register and answer a demographic and expertise survey 325 (LINK). The survey collects information on the analysts current position and 326 self-estimated breadth and level of statistical expertise. We will then provide teams with 327 the acoustic data set and request that they answer the following research question: 328 Do speakers acoustically modify utterances to signal atypical word combinations? Once their analysis is complete, they will answer a structured survey (LINK), 330 providing analysis technique, explanations of their analytical choices, quantitative results, 331 and a statement describing their conclusions. They will also upload their analysis files 332 (including the additionally derived data and text files that were used to extract and 333 pre-process the acoustic data), their analysis code (if applicable), and a detailed 334

36 2.3 Step 3: Peer Reviews of Analyses

journal-ready statistical methods section.

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At a minimum, each analysis will be evaluated by four different reviewers, and each volunteer peer-reviewer will be randomly assigned to methods sections from at least four analyst teams (the exact number will depend on the number of analysis teams and peer reviewers recruited). Each peer reviewer will register and answer a demographic and expertise survey identical to that asked of the analysts. Reviewers will evaluate the methods of each of their assigned analyses one at a time in a sequence determined by the initiating authors. The sequences will be systematically assigned so that, if possible, each analysis is allocated to each position in the sequence for at least one reviewer. For instance, if each reviewer is assigned four analyses to review, then each analysis will be the first

analysis assigned to at least one reviewer, the second analysis assigned to another reviewer, 346 the third analysis assigned to yet another reviewer, and the fourth analysis assigned to a 347 fourth reviewer. Balancing the order in which reviewers see the analyses controls for order 348 effects, e.g. a reviewer might be less critical of the first methods section they read than the 349 last. The process for a single reviewer will be as follows. First, the reviewer will receive a 350 description of the methods of a single analysis. This will include the narrative methods 351 section, the analysis team's answers to our survey questions regarding their methods. 352 including analysis code, and the data set. The reviewer will then be asked, in an online 353 survey (LINK), to rate both the acoustic analysis and the statistical analysis on a scale of 354 0-100 based on these prompts: 355

"Rate the overall appropriateness of the acoustic analysis to answer the research question with the available data. To help you calibrate your rating, please consider the following guidelines:

- 100. A perfect analysis with no conceivable improvements from the reviewer.
- 75. An imperfect analysis but the needed changes are unlikely to dramatically alter final interpretation.
- 50. A flawed analysis likely to produce either an unreliable estimate of the relationship or an over-precise estimate of uncertainty.
- 25. A flawed analysis likely to produce an unreliable estimate of the relationship
 and an over-precise estimate of uncertainty.
- 0. A dangerously misleading analysis, certain to produce both an estimate that is
 wrong and a substantially over-precise estimate of uncertainty that places undue
 confidence in the incorrect estimate.
- *Please note that these values are meant to calibrate your ratings. We welcome ratings of any number between 0 and 100."

After providing this rating, the reviewer will then be shown a series of text boxes and the following prompts:

"Please explain your ratings of this analysis. Please evaluate the selection of acoustic 373 features. Please evaluate the measurement of acoustic features. Please evaluate the choice 374 of statistical analysis type. Please evaluate the process of choosing variables and 375 structuring of the statistical model. Please evaluate the suitability of the variables included 376 in (or excluded from) the statistical model. Please evaluate the suitability of the structure 377 of the statistical model. Please evaluate choices to exclude or not exclude subsets of the 378 data. Please evaluate any choices to transform data (or, if there were no transformations, 379 but you think there should have been, please discuss that choice)." 380

After submitting this review, a methods section from a second analysis will then be 381 made available to the reviewer. This same sequence will be followed until all analyses 382 allocated to a given reviewer have been provided and reviewed. After providing the final 383 review, the reviewer will be simultaneously presented with all four (or more) methods 384 sections that reviewer has just completed reviewing, the option to revise their original 385 ratings, and a text box to provide an explanation. The invitation to revise the original 386 ratings will be as follows: "If, now that you have seen all the analyses you are reviewing, 387 you wish to revise your ratings of any of these analyses, you may do so now." The text box will be prefaced with this prompt: "Please explain your choice to revise (or not to revise) 389 vour ratings." 390

391 2.4 Step 4: Evaluate Variation

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Th initiating authors (SC, JC, TR) will conduct the analyses outlined in this section.

2.4.1 Descriptive statistics. We will calculate summary statistics describing
variation among analyses, including (a) the nature and number of acoustic measures
(e.g. f0 or duration), (b) the operationalization and the temporal domain of measurement

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(e.g. mean of an interval or value at specified point in time), (c) the nature and number of 396 model parameters for both fixed and random effects [if applicable], (d) the nature and 397 reasoning behind inferential assessments (e.g. dichotomous decision based on p-values, 398 ordinal decision based on Bayes factor), as well as the (e) mean, (f) standard deviation and 399 (g) range of the reported effect sizes. 400

Meta-analytical estimation. To summarize the variability in reported 2.4.2401 effect sized, we will follow Bayesian random-effects meta-analytical techniques. Based on 402 the common practices currently in place within the field, we anticipate that researchers will 403 use multi-level/hierarchical/random-effects regression models, so that common effect size 404 measures such as Cohen's d would be inappropriate. Since the variables used by the 405 analysis teams might substantially differ in their measurement scales (e.g., Hz for frequency 406 vs ms for duration), we will standardize all reported effects by refitting each reported 407 model with centered and scaled continuous variables (z-scores, i.e. the observed values 408 subtracted from the mean divided by the standard deviation) and sum-coded factor 409 variables. Factor-level ordering for each factor variable will be decided on a 410 nce good? 411 model-by-model basis, depending on which levels were compared by the team. Each standardized (refitted) model will be fitted as a Bayesian regression model with Stan (Stan 412 Development Team, 2021), RStan (Stan Development Team, 2020), and brms (Bürkner, 2017) in R (R Core Team, 2020). For those reported models that were originally fitted 414 within a frequentist approach, uniform distributions will be used as the priors of all 415 parameters (with the restriction that only positive numbers will be included for scale 416 parameters), making the standardized models in fact equivalent to the reported frequentist 417 models. If a team has fitted Bayesian models, the same priors as reported by the team will 418 be used in fitting the respective standardized model. 419

The estimated coefficients of the critical predictors (i.e. critical according to the 420 analysis teams' self-reported inferential criteria), as obtained from the standardized 421 models, will be used as the standardized effect size (η_i) of each reported model. If multiple 422

predictors within a single analysis have been reported as critical, each will be included in 423 the meta-analytical model (described in details in the next paragraph). Moreover, to 424 account for the differing degree of uncertainty around each standardized effect size, we will 425 use the standard deviation of each effect size returned by the standardized models as the 426 standard error (se_i) of the effect size. This will enable us to fit a so-called 427 "measurement-error" model, in which both the standardized effect sizes and their 428 respective standard errors are entered in the the meta-analytical model. As a desired 429 consequence, effect sizes with a greater standard error will be weighted less than those with 430 a smaller standard error in the meta-analytical calculations. 431

After having obtained the standardized effect sizes η_i with related standard errors se_i, for each critical predictor of the individual reported analyses, the initiating authors will fit a Bayesian random-effects meta-analysis using the following multilevel (intercept-only) regression model:

$$\begin{split} \eta_i &\sim \text{Normal}(\mu_i, \sigma_i) \\ \mu_i &= \alpha + \alpha_{\mathbf{t}[i]} \\ \alpha &\sim \text{Normal}(0, 1) \\ \sigma_{\alpha_\mathbf{t}} &\sim \text{HalfCauchy}(0, 1) \\ \sigma_i &= \mathbf{se}_i \end{split}$$

The outcome variable will be the set of standardized effect sizes η_i . The likelihood of η_i is assumed to correspond to a normal distribution (Knight, 2000). The analysis teams will constitute the group-level effect (i.e., random effect, (1 | team)). The standard errors se_i will be included as the standard deviation σ_i of η_i to fit a measurement-error model, as discussed above. We will use regularizing weakly-informative priors for the intercept α (Normal(0,1)) and for the group-level standard deviation $\alpha_{t[i]}$ (HalfCauchy(0,1)). We will fit this model with 4 chains of Hamiltonian Monte-Carlo sampling for the estimation of

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the joint posterior distribution, using the No U-Turn Sampler (NUTS) as implemented in
Stan, and 4000 iterations (2000 for warm-up) per chain, distributed across 4 processing
This ok?45 cores. In case of issues from divergent transitions in the sampling, we will increase
adapt_delta, tree_depth, and the number of iterations in this order until there are no
divergent transitions. The analysis will be conducted in R (R Core Team, 2020) and fit
using Stan (Stan Development Team, 2021), RStan (Stan Development Team, 2020), and
brms (Bürkner, 2017). The code used to run the model can be found here: INSERT LINK.

The posterior probability of the population-level intercept α will give us an estimate of the range of probable values of the standardized effect size $\hat{\eta}$. This posterior probability will also form the basis of the investigation of the effect of a set of analytical and demographic factors, detailed in Section 2.4.3. Crucially, the posterior probability of σ_{α_t} (the standard deviation of the the group-level effect of team) will allow us to quantify the degree of variation between teams on a standardized scale.

Finally, we will assess whether the standardized effect sizes show bias, and, if so, 456 whether the bias is positive or negative (i.e., whether there is a disproportional greater number of bigger or smaller effect sizes than the meta-analytical mean estimate). This will be achieved through inspection of funnel plots (Light & Pillemer, 1984; for a review see Egger, Smith, Schneider, & Minder, 1997; and Sterne, Becker, & Egger, 2005; for a critique Lau, Ioannidis, Terrin, Schmid, & Olkin, 2006). In brief, a funnel plot is a scatter plot of 461 each standardized effect size with effect size on the x-axis and estimated error 462 (i.e. standard deviation) on y-axis. In absence of bias, the points should be symmetrically 463 distributed around the meta-analytical mean (see Figure 2). A sign of possible bias is when 464 there are more points which are farther from the meta-analytical mean on just one side. 465

2.4.3 Analytical and demographic factors affecting effect sizes. As a second step, we will investigate the extent to which deviations from the meta-analytical mean by individual standardized effect sizes relate to a series of predictors related to analytical and demographic factors (see below). We will fit a single Bayesian regression

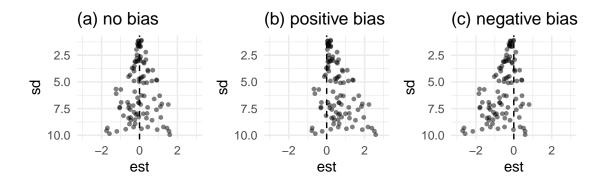


Figure 2. Illustrations of funnel plots showing (a) no bias, (b) positive bias, and (c) negative bias in effect sizes. The vertical dashed line represents the meta-analitical mean.

model with each team's group-level coefficient $\alpha_{\mathbf{t}[i]}$ as the outcome variable, and the analytical and demographic factors described below as predictors. The coefficient $\alpha_{\mathbf{t}[i]}$ can be interpreted as the deviation score of the team's η_i from the meta-analytical mean, allowing us to determine potential effects of analytical and demographic factors on how the teams deviate from the meta-analytical mean. Since each $\alpha_{\mathbf{t}[i]}$ comes with uncertainty, quantified by its standard deviation $\sigma_{\alpha_{\mathbf{t}[i]}}$, we will fit a measurement-error model, as we did in the meta-analytical model above. The formula of the model is the following (see the description of the predictors below for the meaning of acronyms):

$$\begin{split} \alpha_{\mathbf{t}[i]} &\sim \mathrm{Normal}(\mu_{\alpha_{t[i]}}, \sigma_i) \\ \mu_{\alpha_{t[i]}} &= \iota + \upsilon_u \cdot uniq_i + \upsilon_c \cdot cons_i + \upsilon_p \cdot phoc_i + \upsilon_m \cdot modn_i + \upsilon_d \cdot mdim_i + \upsilon_w \cdot twin_i + \upsilon_e \cdot excl_i \\ &\quad + \upsilon_{xp} \cdot rexp_i + \upsilon_{be} \cdot belf_i \\ &\quad \iota \sim \mathrm{Normal}(0, 1) \\ \upsilon_{[\dots]} &\sim \mathrm{Normal}(0, 1) \\ resp_i &\sim \mathrm{Normal}(\mu_{resp_i}, \sigma_{resp_i}) \\ belf_i &\sim \mathrm{Normal}(\mu_{belf_i}, \sigma_{belf_i}) \\ &\quad \sigma_i &= \sigma_{\alpha_{t[i]}} \end{split}$$

The likelihood of $\alpha_{t[i]}$ is a normal distribution. The mean is based on the overall intercept i and on the v coefficient of each predictor. The numeric predictors will be centered and scaled and the categorical predictors will be sum coded. As the prior for i and v we will use a normal distribution with mean 0 and standard deviation 1.

Analytical factors. We will model the effect of the following predictors related to
the analytical characteristics of each team's reported analysis:

- Measure of uniqueness [numeric] of individual analyses for the set of predictors in each model $(uniq_i)$.
- Measure of conservativeness [numeric] of the model specification, as the number of random/group-level effects included $(cons_i)$.
- Number of post-hoc changes to the acoustic measurements [numeric] the teams will report to have carried out $(phoc_i)$.
- Number of models [numeric] the teams will report to have run $(modn_i)$.
- Major dimension [categorical] that has been measured to answer the research question $(mdim_i)$.
- Temporal window [categorical] that the measurement is taken over $(twin_i)$.
- Data exclusion, [categorical] whether data has been excluded or not $(excl_i)$.

The measure of uniqueness of predictors will be assessed by the Sørensen-Dice Index (SDI, Dice, 1945; Sørensen, 1948). The SDI is an index typically used in ecology research to compare species composition across sites. For our purposes, we will treat predictors as species and individual analyses as sites. For each pair of analyses (X, Y), the SDI will be obtained using the following formula:

$$SDI = \frac{2|X \cap Y|}{|X| + |Y|}$$

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where $|X \cap Y|$ is the number of variables common to both models in the pair, and |X| + |Y| is the sum of the number of variables that occur in each model.

In order to generate a unique SDI for each analysis team, we will calculate the
average of all pairwise SDIs for all pairs of analyses using the beta.pair() function in the
betapart R package (Baselga et al., 2020).

The major measurement dimension of each analysis will be categorized according to
the following possible groups: duration, amplitude, fundamental frequency, other spectral
properties (e.g. frequency center of gravity, harmonics difference, etc.), other measures
(e.g. principal components, vowel dispersion, etc.) The temporal window that the
measurement is taken over is defined by the target linguistic unit. We assume the following
relevant linguistic units: segment, syllable, word, phrase.

Demographic factors. We will include the following demographic factors about the teams:

- Research experience [numeric] as the elapsed time from PhD award. Negative values will indicate that the person is a student or graduate studens $(rexp_i)$.
- Initial belief [numeric] in the presence of an effect of atypical noun-adjective pairs on acoustics, as answered during the intake questionnaire $(belf_i)$.

We will publicly archive all relevant data, code, and materials on the Open Science 517 Framework (https://osf.io/3bmcp/). Archived data will include the original data sets 518 distributed to all analysts, any edited versions of the data analyzed by individual groups, 519 and the data we analyze with our meta-analyses, which include the effect sizes derived from separate analyses, the statistics describing variation in model structure among analysis teams, and the anonymized answers to our surveys of analysts and peer reviewers. 522 Similarly, we will archive both the analysis code used for each individual analysis and the 523 code from our meta-analyses. We will also archive copies of our survey instruments from 524 analysts and peer reviewers. 525

Our rules for excluding data from our study are as follows. We will exclude from our synthesis any individual analysis submitted after we have completed peer review or those unaccompanied by analysis files that allow us to understand what the analysts did. We will also exclude any individual analysis that does not produce an outcome that can be interpreted as an answer to our primary question.

2.5 Step 6: Collaborative Write-Up of Manuscript

Analysts and initiating authors will discuss the limitations, results, and implications
of the study and collaborate on writing the final manuscript for review as a stage-2
Registered Report.

3 References

- Arts, A., Maes, A., Noordman, L. G., & Jansen, C. (2011). Overspecification in written instruction. *Linguistics*, 49(3), 555–574.
- Baselga, A., Orme, D., Villeger, S., De Bortoli, J., Leprieur, F., & Logez, M. (2020).

 betapart: Partitioning beta diversity into turnover and nestedness components.

 Retrieved from https://CRAN.R-project.org/package=betapart
- Botvinik-Nezer, R., Holzmeister, F., Camerer, C. F., Dreber, A., Huber, J.,

 Johannesson, M., ... others. (2020). Variability in the analysis of a single

 neuroimaging dataset by many teams. *Nature*, 1–7.
- Box, G. E. (1976). Science and statistics. Journal of the American Statistical

 Association, 71(356), 791–799.
- Brugger, P. (2001). From haunted brain to haunted science: A cognitive

 neuroscience view of paranormal and pseudoscientific thought. *Hauntings and*Poltergeists: Multidisciplinary Perspectives, 195–213.
- Bürkner, P.-C. (2017). brms: An R package for Bayesian multilevel models using

 Stan. Journal of Statistical Software, 80(1), 1–28.

 https://doi.org/10.18637/jss.v080.i01
- Camerer, C. F., Dreber, A., Holzmeister, F., Ho, T.-H., Huber, J., Johannesson, M.,

 ... others. (2018). Evaluating the replicability of social science experiments in

 nature and science between 2010 and 2015. Nature Human Behaviour, 2(9),

 637–644. https://doi.org/10.1038/s41562-018-0399-z
- De Groot, A. D. (2014). *Thought and choice in chess* (Vol. 4). Walter de Gruyter
 GmbH & Co KG.
- Degen, J., Hawkins, R. D., Graf, C., Kreiss, E., & Goodman, N. D. (2020). When redundancy is useful: A bayesian approach to "overinformative" referring

- expressions. Psychological Review.
- Dice, L. R. (1945). Measures of the amount of ecologic association between species.

 Ecology, 26(3), 297–302.
- Egger, M., Smith, G. D., Schneider, M., & Minder, C. (1997). Bias in meta-analysis detected by a simple, graphical test. *British Medical Journal*, 315, 629–634.
- Fischhoff, B. (1975). Hindsight is not equal to foresight: The effect of outcome knowledge on judgment under uncertainty. *Journal of Experimental Psychology:*Human Perception and Performance, 1(3), 288.
- Flake, J. K., & Fried, E. I. (2019). Measurement schmeasurement: Questionable
 measurement practices and how to avoid them. Pre-print available at PsyArXiv.
- Foulkes, P., & Docherty, G. (2006). The social life of phonetics and phonology.

 Journal of Phonetics, 34(4), 409–438.
- Gatt, A., Gompel, R. P. van, Deemter, K. van, & Krahmer, E. (2013). Are we bayesian referring expression generators. Cognitive Science Society.
- Gelman, A., & Loken, E. (2014). The statistical crisis in science: Data-dependent
 analysis—a" garden of forking paths"—explains why many statistically significant
 comparisons don't hold up. American Scientist, 102(6), 460–466.
- Gordon, M., & Roettger, T. (2017). Acoustic correlates of word stress: A

 cross-linguistic survey. *Linguistics Vanguard*, 3(1), 1–11.
- Grice, H. P. (1975). Logic and conversation. In Speech acts (pp. 41–58). Brill.
- Hawkins, S., & Nguyen, N. (2004). Influence of syllable-coda voicing on the acoustic properties of syllable-onset /l/ in English. *Journal of Phonetics*, 32(2), 199–231.
- Jongman, A., Wayland, R., & Wong, S. (2000). Acoustic characteristics of english fricatives. The Journal of the Acoustical Society of America, 108(3), 1252–1263.
- Knight, K. (2000). Mathematical statistics. Chapman; Hall, New York.

- Koole, S. L., & Lakens, D. (2012). Rewarding replications: A sure and simple way to improve psychological science. *Perspectives on Psychological Science*, 7(6), 608–614.
- Ladd, D. R. (2008). *Intonational phonology*. Cambridge University Press.
- Lau, J., Ioannidis, J. P. A., Terrin, N., Schmid, C. H., & Olkin, I. (2006). The case of the misleading funnel plot. *British Medical Journal*, 333(7568), 597–600.
- Light, R. J., & Pillemer, D. B. (1984). Summing up; the science of reviewing

 research. Cambridge, MA (USA) Harvard Univ. Press.
- Lisker, L. (1977). Rapid versus rabid: A catalogue of acoustic features that may cue
 the distinction. The Journal of the Acoustical Society of America, 62(S1),
 S77–S78.
- Lisker, L. (1986). "Voicing" in English: A catalogue of acoustic features signaling

 /b/ versus /p/ in trochees. Language and Speech, 29(1), 3–11.
- Nickerson, R. S. (1998). Confirmation bias: A ubiquitous phenomenon in many guises. Review of General Psychology, 2(2), 175–220.
- Nosek, B. A., & Lakens, D. (2014). A method to increase the credibility of published results. *Social Psychology*, 45(3), 137–141.
- Ogden, R. (2004). Non-modal voice quality and turn-taking in Finnish. Sound

 Patterns in Interaction: Cross-Linguistic Studies from Conversation, 29–62.
- Open Science Collaboration. (2015). Estimating the reproducibility of psychological science. Science, 349(6251). https://doi.org/10.1126/science.aac4716
- Paraboni, I., Van Deemter, K., & Masthoff, J. (2007). Generating referring

 expressions: Making referents easy to identify. Computational Linguistics, 33(2),

 229–254.

- R Core Team. (2020). R: A language and environment for statistical computing.

 Vienna, Austria: R Foundation for Statistical Computing. Retrieved from

 https://www.R-project.org/
- Roettger, T. B. (2019). Researcher degrees of freedom in phonetic research.

 Laboratory Phonology: Journal of the Association for Laboratory Phonology,

 10(1).
- Rotello, C. M., Heit, E., & Dubé, C. (2015). When more data steer us wrong:

 Replications with the wrong dependent measure perpetuate erroneous

 conclusions. *Psychonomic Bulletin & Review*, 22(4), 944–954.
- Rubio-Fernández, P. (2016). How redundant are redundant color adjectives? An
 efficiency-based analysis of color overspecification. Frontiers in Psychology, 7,

 153.
- Sedivy, J. C. (2003). Pragmatic versus form-based accounts of referential contrast:

 Evidence for effects of informativity expectations. *Journal of Psycholinguistic*Research, 32(1), 3–23.
- Silberzahn, R., Uhlmann, E. L., Martin, D. P., Anselmi, P., Aust, F., Awtrey, E., ...
 others. (2018). Many analysts, one data set: Making transparent how variations
 in analytic choices affect results. Advances in Methods and Practices in

 Psychological Science, 1(3), 337–356.
- Simmons, J. P., Nelson, L. D., & Simonsohn, U. (2011). False-positive psychology:

 Undisclosed flexibility in data collection and analysis allows presenting anything
 as significant. *Psychological Science*, 22(11), 1359–1366.
- Stan Development Team. (2020). RStan: The R interface to Stan. Retrieved from http://mc-stan.org/

647

649

- Stan Development Team. (2021). Stan modeling language users guide and reference manual, v2.26.0. Retrieved from http://mc-stan.org/
- Starns, J. J., Cataldo, A. M., Rotello, C. M., Annis, J., Aschenbrenner, A., Bröder,
 A., ... Wilson, J. (2019). Assessing theoretical conclusions with blinded inference
 to investigate a potential inference crisis. Advances in Methods and Practices in

 Psychological Science, 2(4), 335–349.
- Sterling, T. D. (1959). Publication decisions and their possible effects on inferences
 drawn from tests of significance—or vice versa. *Journal of the American*Statistical Association, 54(285), 30–34.
- Sterne, J. A. C., Becker, B. J., & Egger, M. (2005). The funnel plot. In H. R.

 Rothstein, A. J. Sutton, & M. Borenstein (Eds.), *Publication bias in*meta-analysis: Prevention, assessment and adjustments (pp. 75–98). Wiley

 Online Library.
 - Sørensen, T. (1948). A method of establishing groups of equal amplitude in plant sociology based on similarity of species and its application to analyses of the vegetation on Danish commons. Kongelige Danske Videnskabernes Selskab, 5(4), 1–34.
- Tukey, J. W. (1977). Exploratory data analysis (Vol. 2). Reading, MA.
- Tversky, A., & Kahneman, D. (1974). Judgment under uncertainty: Heuristics and biases. *Science*, 185(4157), 1124–1131.
- Van Heuven, V. J., Haan, J., Gussenhoven, C., & Warner, N. (2002). Temporal distribution of interrogativity markers in Dutch: A perceptual study. In

 Laboratory phonology 7 (Vol. 4, p. 61). Walter de Gruyter.
- Van Heuven, V. J., & Van Zanten, E. (2005). Speech rate as a secondary prosodic characteristic of polarity questions in three languages. *Speech Communication*,

47(1-2), 87-99.

- Wagenmakers, E.-J., Wetzels, R., Borsboom, D., Maas, H. L. van der, & Kievit, R.

 A. (2012). An agenda for purely confirmatory research. *Perspectives on*Psychological Science, 7(6), 632–638.
- Westerbeek, H., Koolen, R., & Maes, A. (2015). Stored object knowledge and the production of referring expressions: The case of color typicality. Frontiers in Psychology, 6, 935.
- White, L., Payne, E., & Mattys, S. L. (2009). Rhythmic and prosodic contrast in
 Venetan and Sicilian Italian. In M. Vigario, S. Frota, & M. J. Freitas (Eds.),

 Phonetics and phonology: Interactions and interrelations (pp. 137–158).

 Amsterdam: John Benjamins.
- Wicherts, J. M., Borsboom, D., Kats, J., & Molenaar, D. (2006). The poor
 availability of psychological research data for reanalysis. *American Psychologist*,
 61(7), 726.
- Winter, B. (2014). Spoken language achieves robustness and evolvability by exploiting degeneracy and neutrality. *BioEssays*, 36(10), 960–967.