The Unreliability of Measures of Intercoder Reliability, and What to do About it

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Abstract

In both automated and traditional text analysis, human coders are regularly tasked with categorizing documents. Researchers then evaluate the success of this crucial step in the research process via one of many measures of intercoder reliability, such as Cronbachs alpha. They then improve coding practices until this measure reaches some arbitrary threshold, at which point remaining disagreements are resolved in arbitrary ways and ignored in subsequent analyses. We show that this common practice can generate severely biased estimates and misleading conclusions. The problem is the focus on measures of intercoder reliability which, except at the extreme, are unrelated to the quantities of interest, such as the proportion of documents in each category. We thus develop an approach that enables scholars to directly incorporate coding uncertainty into statistical estimation. The method offers an interval estimate which we prove contains the true proportion of documents in each category, under reasonable assumptions. We then extend this method to situations with multiple coders, when one coder is trusted more than another, and when the resulting document codes are used as inputs to another statistical model. We offer easy-to-use software that implements all our suggestions.

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1   Introduction

We attempt to shore up an essential and well known, but relatively undiscussed, component of the science of automated and traditional text analysis. Researchers in these areas spend considerable time and resources managing teams that code documents into a chosen set of categories. Categorization is a fundamental component of human understanding, but the ambiguous and ever creative nature of human language makes the process difficult and perfection impossible (**?**, **?**, **?**, Appendix, **?**, p.10). The consequences for the ultimate conclusions drawn depend crucially on the extent and nature of these difficulties in this formative stage of analysis.

The usual practice involves defining a set of mutually exclusive and exhaustive categories based on substantive interest and a theoretical perspective; assigning two or more research assistants to code a small set of documents into these categories; discovering that levels of intercoder reliability are too low; adjusting, redefining, and clarifying the categories; and then retraining the research assistants and coding a new set of documents from scratch. This process is repeated until the level of intercoder reliability is deemed satisfactory. Although the level of intercoder reliability reached often goes unreported in many fields Lombard, Snyder-Duch and Bracken (2002); Riffe and Freitag (1997), the threshold "satisfactory" level in most fields is approximately 70–80% (**?**, pp. 4–5). At that point, the disagreements are ignored. The remaining documents are coded by a single coder, or by multiple coders who negotiate their differences informally or their codes are averaged. This practice is common across fields, including political science (Stewart and Zhukov, 2009; De Vreese et al., 2006; Druckman and Parkin, 2005; Jamal et al., 2014; Druckman, Kifer and Parkin, 2010, 2009), medicine (Hripcsak and Heitjan, 2002), education (Béchard and Grégoire, 2005), journalism (Leccese, 2009), sociology (Williams et al., 2009), business and marketing (Nazli Nik Ahmad and Sulaiman, 2004; Bortree and Seltzer, 2009), psychology (Zullow et al., 1988), and communication (Van Gorp, 2005).

For traditional hand coding projects, all documents are coded in this way. For many types of automated text analysis, only documents in the training set are coded via these procedures. For other types of machine learning analyses, the training set has high accuracy (due to known information, such as the author of a speech), but the classifier used to sort the test set into categories produces analogous errors that are typically as large as hand coding exercises. For any of these, the consequences for ignoring the remaining errors be devastating for quantities of interest of special interest to social scientists. In particular, social scientists usually have little interest in the numeric code for any one social media post or congressional speech (which they could of course merely read). Instead, the quantity of interest in the social sciences usually involves the percent of documents in each category, such as treaties involving strategic misperception or social media posts supporting each presidential candidate.

Our proposal to ameliorate this problem does not involve changing the best practices of coders in improving their levels of intercoder reliability. Researchers should continue the practice of improving their categories and training their coders, as has been the case (e.g., **?**). We also have no objection to the long history of work on measures of intercoder reliability, such as Cronbach's α (Cronbach, 1951), Scott's

π (Scott, 1955), Cohen's κ (Byrt, Bishop and Carlin, 1993), Krippendorff's α (Hayes and Krippendorff, 2007), and many others. This literature continues to provide more nuanced, improving ways of summarizing a confusion matrix in a scalar metric for intercoder reliability. This work should also continue.

The problem then is neither the researcher's coding practices nor the methodologist's measures of intercoder reliability. The problem is the near universal practice of using these measures to determine a threshold level, above which intercoder reliability may be safely ignored. In fact, as we show, no such level exists. Other than near perfect agreement, which does not occur with meaningful categories and documents, no level of intercoder reliability should be ignored. What is missing then, is a set of tools that enable researchers to incorporate the remaining disagreement in their analytical methods. We provide those here. Although generic methods of measurement error have been proposed (**?**, Appendix; **?**, Molinari 2008), the methods introduced here are tuned for exactly this problem and so are potentially more powerful with fewer assumptions.

2 The Problem with Using Reliability to Assess Validity

In this section, we derive a mathematical relationship between reliability to validity. For now, we assume the absence of sampling variability, and use "probability" and "proportion" interchangably. We begin with notation, which is also summarized as Appendix .

Consider C coders ($c = 1, \ldots, C$) tasked with coding D_i ($d = 1, \ldots, D_i$) documents into K ($k = 1, \ldots, K$) categories. Each document d has a true category $\pi_d \in \{1, \ldots, K\}$, and each category k has a true proportion of documents, $\bar{\pi}_k = \text{mean}_d[I(\pi_d = k)]$, which we collect as a vector $\bar{\boldsymbol{\pi}} = \{\bar{\pi}_i : k = 1, \ldots, K\}$ (and where $\text{mean}_j(a_j) = \sum_{j=1}^{J} a_j / J$ and J is the number of objects contributing to the summation). Analogously, each coder c's decision on document d is $y_d^c \in \{1, \ldots, K\}$ and the proportion of documents coder c puts in category k is $\bar{y}_k^c = \text{mean}_d[I(y_d^c = k)]$. We also collect these results in a vector, $\bar{\boldsymbol{y}}^c = (\bar{y}_1^c, \bar{y}_2^c, \ldots, \bar{y}_K^c)$. For a raw or naive estimate, we average these results over coders, producing $\bar{y}_k = \text{mean}_c(\bar{y}_k^c)$ with vector $\bar{\boldsymbol{y}} = (\bar{y}_1, \bar{y}_2, \ldots, \bar{y}_K)$.

We now explicitly relate estimates to the truth by explicitly writing the "data coding generation process". We begin in scalar notation with the observed proportion in category k as coded by c as a function of all the true proportions:

$$\bar{y}_k^c = \sum_{j=1}^{K} \epsilon_{jk}^c \bar{\pi}_j$$

where ϵ_{jk}^c is misclassification probability — the probability that coder c classifies a document into category k if it is in fact in category j. This means that $\sum_{j=1}^{J} \epsilon_{jk}^c = 1$ because the coder must make some decision with each document. We define ϵ_{kk}^c as coder c's *validity* for category k: the proportion of documents the coder correctly classifies in category k from category k, so that if $\epsilon_{kk}^c = 1$, then $\bar{y}_k^c = \bar{\pi}_k$.

We then move to matrix notation by collecting ϵ_{jk}^c misclassification probabilities into a $K \times K$ misclassification matrix $\boldsymbol{E}^c$, with coder c's validities ϵ_{kk}^c on the diagonal. The off diagonal entries measure the probabilities of errors that coder c makes. We then write the observed proportions for coder c simply as (see Kuha and Skinner, 1997; Mikhaylov, Laver and Benoit, 2012),

$$\bar{\boldsymbol{y}}^c = \boldsymbol{E}^c \bar{\boldsymbol{\pi}} \tag{2.1}$$

If we observe $\boldsymbol{E}^c$ then, a simple correction reveals the true $\boldsymbol{\pi}$,

$$(\boldsymbol{E}^c)^{-1} \bar{\boldsymbol{y}}^c = \bar{\boldsymbol{\pi}} \tag{2.2}$$

We will represent the proportion of the time coder c is correct as $\epsilon^c = \sum_{k=1}^K \epsilon_{kk}^c \bar{\pi}_k$. Note, this can also be interpreted as the $\bar{\boldsymbol{\pi}}$ weighted average validity.

Of course not only is $\boldsymbol{E}^c$ rarely if ever observed, but researchers usually do not include features designed to estimate or influence it. Thus, instead of focusing on *validity*, researchers are often encouraged to focus on the agreement between coders, which we refer to as *reliability*. To do this, researchers set up coding tasks so that at least two coders categorize a subset of documents. The intuition expressed in the literature is that improving reliability may also improve validity, although the feeling is not universal (**?**, p. 130). A key goal of this paper is to formalize these intuitions and connect these two concepts mathematically.

If two coders, 1 and 2, both code D documents, we define a confusion matrix element as the probability that coder 1 classifies a document as j and coder 2 classifies a document as k as $m_{jk}^{12} = \text{mean}_d[I(y_d^1 = j, y_d^2 = k)]$. The category-specific reliabilities are on the diagonal of this matrix, m_{kk}^{12}. Finally, as a summary, we define the overall reliability, the proportion of times coders 1 and 2 apply the same labels across all categories, as the sum of the diagonal elements of the confusion matrix:

$$a^{12} = \sum_{k=1}^K m_{kk}^{12}$$

Some of the alternative scalar measures of reliability often begin with a^{12} and adjust for some concept of base levels of agreement that could be achieved by chance. The differences among these measures can be important for some purposes, but they do not materially affect our results.

Regardless of the measure used, researchers are often advised to focus on improving reliability, which is readily measured. Then, once reliability is improved as much as feasible given resources and time constraints, coder discrepancy is usually ignored for subsequent analyses. We show the problem with this approach by demonstrating that even high (but not perfect) levels of reliability do not satisfactorily constrain validity.

We begin formally connecting validity and reliability with Proportion 1

Proposition 1. *1) Suppose that $K > 2$ and coder 1 and coder 2 have overall reliability a^{12}. Then coder 1's average validity is $\epsilon^1 \in [0, 1]$ and coder 2's validity is $\epsilon^2 \in [0, 1]$.*

4

2) *Suppose $K \geq 2$. Then coder 1 and coder 2 have maximum average validity $\dot{\epsilon}^{12} = \frac{1+a^{12}}{2}$ and if the coders have the maximum average validity $\epsilon^1 = a^{12} + 1 - \epsilon^2$.*

Proof. Consider first the individual validity intervals for the coder. Fix an agreement rate between the coders as a^{12}. Suppose the coders agree on the same mistake a^{12} of the time and disagree and make distinct errors $1 - a^{12}$ of the time. Then if $K > 2$, $\epsilon^1 = 0$, $\epsilon^2 = 0$. If coder 1 is always correct and coder 2 is correct a^{12}, then $\epsilon^1 = 1$. If coder 2 is always correct and coder 1 is correct a_{12} then $\epsilon^2 = 1$.

Now consider the maximum joint validity $\dot{e}$. Note that for all pairs of documents the coders either agree or they disagree. For the coders to have maximum joint validity given the agreement rate then when they agree the coders must be correct and when they disagree one coder must be correct. Thus $a^{12} = P(\text{Coder 1 and Coder 2 Correct})$ and that $P(\text{Coder 1 Correct or Coder 2 Correct}) = 1$. Then, by inclusion/exclusion theorem:

$$
\begin{aligned}
a^{12} &= P(\text{Coder 1 Correct}) + P(\text{Coder 2 Correct}) - P(\text{Coder 1 Correct or Coder 2 Correct}) \\
&= \epsilon^1 + \epsilon^2 - 1
\end{aligned}
$$

by definition of ϵ^1 and ϵ^2.

Algebraic manipulation then shows that the maximum joint validity $\dot{e} = \frac{1+a^{12}}{2}$ and that at this maximum $\epsilon^1 = a^{12} + 1 - \epsilon^2$.

$\square$

Proposition 1 shows that without additional behavioral assumptions, even high coder agreement does not imply that we know that our coders are performing accurately. Even if coders agree perfectly, it may be the case that they always agree in error. To eliminate this possibility, we need to make an assumption that our coders are not *malicious*: able to perfectly misrepresent the answers in our study. Of course, we are often able to eliminate this assumption with careful selection, training, and monitoring of our coders. But obtaining a lower bound on the joint accuracy would still rely heavily on how pessimistic we are about our coders' joint validity. In Proposition 4 below show that the coders' joint validity can range from zero to the maximum described in Proposition 1. Therefore, an assumption alone would define the lower bound on the coders' performance.

Rather than focus on the pessimistic case, we first make the optimistic assumption that our coders are performing as well as possible, given their level of agreement. Formally, this is equivalent to assuming $\dot{e} = \frac{\epsilon^1 + \epsilon^2}{2}$ is at a maximum, given the coders' reliability. Proposition 1 does show that agreement provides an upper bound on the joint accuracy of our coders. We show that in the next section that assuming our coders have the maximum validity given their codes provides much more informative intervals. We view this assumption as much less optimistic than the current approach—assuming no coder error remains for high levels of agreement—but not so unrealistically pessimistic as to require an assumption that our coders are intentionally coordinating to undermine our project. We will also see that our framework will provide a natural sensitivity analysis to relax this optimistic assumption to the more pessimistic case where our coders are performing more poorly and where coders

make non-constant errors on the off-diagonals. We show that this generalization will correct two instances where the standard parametric models of coder performance tend to perform poorly.

Proposition 1 leads to our first assumption:

Assumption 1. ***Wisdom of the Coders***. *We will suppose that our coders' average validity is at a maximum. Equivalently, as we show below, this means if a plurality of coders of document d agree, they agree on the truth; if there is a tie, then at least one side identifies the truth.*

3   A Method for Incorporating Coder Error

In this section we will develop an algorithm that allows us to propagate the uncertainty from our coders, using Assumptions 2 and 1. Rather than only a point estimate, we take a partial identification approach resulting in a procedure that returns an interval that contains true value if the assumptions are satisfied. We first provide intuition about how to use evaluation matrices to obtain bounds on the true proportions in each category.

3.1   Intuition about Bounds

To gain intuition about our approach and the errors from categorical measurement error consider the simplest coding situation. Suppose that a coder is attempting to classify documents into two categories.

We suppose that our single coder is able to perfectly classify documents that truly belong in category 2, but is only correct ϵ_{11}^1 of the time when a document truly belongs to category 1. We can represent the coder's decisions with the following evaluation matrix

$$\boldsymbol{E}^1 \;\; = \;\; \begin{pmatrix} \epsilon_{11}^1 & 0 \\ (1 - \epsilon_{11}^1) & 1 \end{pmatrix}$$

We can then represent the observed proportions, $\boldsymbol{E}^1 \bar{\boldsymbol{\pi}} = \bar{\boldsymbol{y}}^1 = (\epsilon_{11}^1 \bar{\pi}_1^1, (1 - \epsilon_{11}^1)\bar{\pi}_1 + \bar{\pi}_2)$, where $\boldsymbol{\pi} = (\pi_1, \pi_2)$ is the true proportion in each category.

If we suppose that ϵ_{11}^1 is known, then we can solve for the true values of $\bar{\boldsymbol{\pi}}$. They are

$$\bar{\pi}_1 \;\; = \;\; \frac{\bar{y}_1^1}{\epsilon_{11}^1} \tag{3.1}$$

$$\bar{\pi}_2 \;\; = \;\; \bar{y}_2^1 - \frac{1 - \epsilon_{11}^1}{\epsilon_{11}^1}\bar{y}_1^1 \tag{3.2}$$

Both equations are intuitive. The first notes that the only way our coder would classify something in category 1 is if it truly belongs to category 1, because she perfectly handles all documents that correctly belong to category 2. The proportion in Category 1 will be too small and dividing by ϵ^1 correctly adjusts the proportion.

The second equation adjusts $\bar{y}_2^1$, removing $(1-\epsilon^1)$ of the incorrectly coded proportion of documents that should have been coded in Category 1 $\frac{\bar{y}_1^1}{\epsilon^1}$.

Suppose, now, that we don't know that the true values of ϵ^1, but rather some interval in which the values might lie—say $\epsilon^1 \in [a, 1]$. Then, we can plug in the values in Equation 3.1 and 3.2 to determine bounds on the proportion in each category. In particular, for this interval we find that $\pi_1 \in [\frac{\bar{y}_1^1}{1}, \frac{\bar{y}_1^1}{a}]$ and $\pi_2 \in [\bar{y}_2^1 - \frac{1-a}{a}\bar{y}_1^1, \bar{y}_2^1]$. Unlike other continuous measurement error models, note that the observed proportions define the lower-bound for Category 1 and the upper bound for Category 2.

Throughout the paper we will consider a more general setting, where coders make classification decisions across several categories. To motivate our first interval estimator we will suppose that our coders make constant errors across categories—an assumption that is useful for motivating intuition and as a default method, but an assumption that we relax below. Specifically we will suppose that $\epsilon^1 = \epsilon_{11}^1 = \epsilon_{22}^1 = \epsilon_{33}^1$.

$$\boldsymbol{E}^1 = \begin{pmatrix} \epsilon^1 & \epsilon_{12}^1 & \epsilon_{13}^1 \\ \epsilon_{21}^1 & \epsilon^1 & \epsilon_{23}^1 \\ \epsilon_{31}^1 & \epsilon_{32}^1 & \epsilon^1 \end{pmatrix}$$

Numerical Example to Obtain Bounds   The reasoning thus far has been about a simple case with one coder. To derive our bounds we will use the information from double coded data to obtain information about each coders' validity and to therefore obtain bounds on the true proportion in each category. For a numerical example, we will suppose that the true proportion in each category is $\bar{\pi} = (0.7, 0.25, 0.05)$ and that the coders' proportions are,

$$\bar{\boldsymbol{y}}^1 = \begin{pmatrix} 0.9 & 0.07 & 0.02 \\ 0.08 & 0.9 & 0.08 \\ 0.02 & 0.03 & 0.9 \end{pmatrix} \bar{\pi} = (0.65, 0.28, 0.07)$$

$$\bar{\boldsymbol{y}}^2 = \begin{pmatrix} 0.8 & 0.14 & 0.17 \\ 0.01 & 0.80 & 0.03 \\ 0.19 & 0.06 & 0.8 \end{pmatrix} \bar{\pi} = (0.6, 0.21, 0.19)$$

With naive estimate $\bar{\boldsymbol{y}} = (0.63, 0.25, 0.13)$. We will suppose that our coders have agreement rate 0.7.

Our approach to obtaining bounds will be to make assumptions that make use of information about our coders' agreement rate and their estimated proportions to inform the bounds that are used on the proportions. To gain intuition, we will walk through a set of assumptions that place increasingly specific assumptions on the validity and therefore narrow the bounds. We will focus on obtaining bounds for the first category.

First, we will merely assume that our coders perform better than some baseline level and not make user of agreement rate. Specifically, we will initially suppose that our coders perform above some lower level of validity, $\epsilon^1 \in [0.65, 1]$ and $\epsilon^2 \in [0.65, 1]$ and make no other assumptions about the errors. As we show in Table 3.1, this provides fairly wide intervals for each of the proportions.

	Category 1	Category 2	Category 3
		Table 1: Intuition About Bounds	
Truth	0.7	0.25	0.05
Naive Estimate	0.63	0.25	0.13
Constant Validity ($\epsilon^1, \epsilon^2 \in [0.65, 1]$)	[0.63, 0.88]	[0.00, 0.29]	[0.00, 0.18]
Maximum Average Validity	[0.68, 0.77]	[0.09, 0.29]	[0.00, 0.16]
MAV + Structure	[0.69, 0.73]	[0.21, 0.26]	[0.02, 0.08]

In what follows we will make the more restrictive assumption that our coders' have maximum average validity, given their agreement rate. We then provide a method for relaxing this assumption and exploring different behavioral assumptions about how our coders are performing. Making the maximum average validity assumption narrows the intervals further. Our method will combine this assumption with particular information about the structure of the problem and that any solution must maintain. The final row in Table 3.1 shows that using these further properties further narrows the interval for each proportion, while ensuring that the truth lies in each interval.

In what follows we use the intuition and assumptions from this section to derive a more general algorithm for estimating bounds on the proportion in categories.

3.2   Deriving the Intervals

To motivate our method for obtaining intervals, recall that Equation 2.2 shows that if we know $\boldsymbol{E}^c$ for coder c then we can use the coder's estimates to back out the true values $\bar{\boldsymbol{\pi}}$. Of course we do not know the true values of $\boldsymbol{E}^c$, but Proposition 1 provides an upper bound on the coders' validity—it defines a range for each coder that depends on the agreement between the pair of coders and the other coder's validity. Using this information, and other properties that a solution must have, we can define a set of pairs of matrices where the coders have maximum average validity. We then find the minimum and maximum values of $\bar{\pi}_k$ over this set of matrices.

To derive this interval, we first make the maximum average validity assumption and to simplify our analysis we first assume our coders have constant accuracy across categories.

Assumption 2. ***Constant Validity Assumption*** *Coder c's validity is constant across categories. This implies that it can be simplified as $\epsilon^c = \epsilon^c_{kk}$ for all k.*

Assumptions 1 and 2 are restrictive, but we show how to directly relax them in the following sections, making less optimistic assumptions about coder accuracy and more general assumptions about the kinds of errors coders make. Note that under Assumption 2 the diagonal of the evaluation matrix for each coder $\boldsymbol{E}^c$ is a constant.

The upper bound from Proposition 1 defines a range of potential pairs of values for ϵ^1 and ϵ^2: $\epsilon^2 \in [a^{12}, 1]$ and $\epsilon^1 = a^{12} + 1 - \epsilon^2$. For intuition note that if $\epsilon^2 = a^{12}$ then $\epsilon^1 = 1$ and if $\epsilon^2 = 1$ then $\epsilon^1 = a^{12}$.

Proposition 1 provides values for the diagonal elements of $\boldsymbol{E}^1, \boldsymbol{E}^2$, but provides no information about the off-diagonal elements, which we will optimize over to obtain intervals for each category. Before performing this optimization, however, we can use further constraints to limit the number of matrices we search over. First, as demonstrated in the previous section, if we know that if we have the true evaluation matrices, then $\left(\boldsymbol{E}^1\right)^{-1}\bar{\boldsymbol{y}}^1 = \left(\boldsymbol{E}^2\right)^{-1}\bar{\boldsymbol{y}}^2$. Second, we know that any solution must have all entries between zero and one. Therefore, we can restrict our attention to pairs of matrices $(\boldsymbol{E}^c)^{-1}\bar{\boldsymbol{y}}^c \in \Delta^{K-1}$, where Δ^{K-1} is the $K-1$ dimensional simplex, for all coders c. Third, because of the structure of the evaluation matrices, we know that each column must sum to 1, so the off-diagonal elements for each column must sum to $1 - \epsilon^c$ for $\boldsymbol{E}^c$. We can collect the pairs of matrices that satisfies these constraints into the set $\mathbb{E}$ with typical element $(\tilde{E}^1, \tilde{E}^2) \in \mathbb{E}$.

We can then optimize over the set of matrix pairs $\mathbb{E}$ to obtain an interval estimator for π_k. Note that because we have restricted $\mathbb{E}$ to contain matrices such that $(\tilde{E}^1, \tilde{E}^2) \in \mathbb{E}$ implies that $\left(\tilde{\boldsymbol{E}}^1\right)^{-1}\bar{\boldsymbol{y}}^1 = \left(\tilde{\boldsymbol{E}}^2\right)^{-1}\bar{\boldsymbol{y}}^2$, we know that

$$\min_{(\tilde{E}^1, \tilde{E}^2) \in \mathbb{E}} \frac{\left(\tilde{\boldsymbol{E}}^1\right)^{-1}\bar{\boldsymbol{y}}^1 + \left(\tilde{\boldsymbol{E}}^2\right)^{-1}\bar{\boldsymbol{y}}^2}{2} = \min_{(\tilde{E}^c) \in \mathbb{E}} \left(\tilde{\boldsymbol{E}}^c\right)^{-1}\bar{\boldsymbol{y}}^c$$

for coders $c = 1, 2$.

We will therefore define our interval estimator for $\bar{\pi}_k$ as $\bar{\pi}_k^{\text{int}}$

$$\bar{\pi}_k^{\text{int}} = \left[\min_{\tilde{E}^c \in \mathbb{E}} \left(\tilde{\boldsymbol{E}}^c\right)^{-1}\bar{\boldsymbol{y}}^c|_k, \max_{\tilde{E}^c \in \mathbb{E}} \left(\tilde{\boldsymbol{E}}^c\right)^{-1}\bar{\boldsymbol{y}}^c|_k\right] \tag{3.3}$$

where $|_k$ denotes selecting the kth element from a vector.

Proposition 2 shows that under our assumption of maximum average validity (Assumption 1) and constant category validity assumption (Assumption 2), $\bar{\pi}_k^{\text{int}}$ will contain the true values of $\bar{\pi}_k$. Further, the bounds are sharp, in the sense that making any additional use of the coding decisions—such as the error structure—would require an additional set of assumptions about the proportions in the categories or further additional assumptions about each coders' validity.

Proposition 2. *Suppose that coder 1 and coder 2 have agreement a^{12} and that Assumptions 1 and 2 hold. Then $\bar{\pi}_k \in \bar{\pi}_k^{int}$ for each k.*

Proof. Assumptions 1 and 2 imply that the true evaluation matrices $(\boldsymbol{E}^1, \boldsymbol{E}^2) \in \mathbb{E}$. This follows because at the true evaluation matrices the coders will have constant validity, by Assumption 2, the inverted evaluation matrices will yield the correct answer and therefore be equal to each other, and the true proportions lie in the simplex. From Equation 2.2 this implies that $(\boldsymbol{E}^c)^{-1}\bar{\boldsymbol{y}}^c = \bar{\boldsymbol{\pi}}$ for both coder $i = 1, 2$. Thus $\bar{\pi}_k \in \bar{\pi}_k^{\text{int}}$. $\square$

To obtain the interval that coder agreement, coders' decisions, and maximum joint validity implies we optimize over the pairs of matrices that are possible under the assumptions to find the minimum and maximum implied values of $\bar{\pi}_k$. The

9

matrix inverses in Equation 3.3 make straightforward optimization difficult to obtain the identification region. Therefore, we use an iterative algorithm to obtain the interval estimates for $\bar{\pi}_k^{\text{int}}$. We describe this algorithm to obtain the interval in Appendix B.

3.3   What If There is No Truth?

Our algorithm is based on the assumption that there is a true proportion of documents in each category. This assumption will hold in many situations: coding rules should be written so that there is an unambiguous true proportion in each category. Yet, some scholars might dispute this assumption is reasonable. Instead they might argue that the truth depends on the model that coders have in mind. Even in the absence of an assumption of a gold standard truth our method still provides a useful and intuitive method for incorporating unceratinty from our hand coders.

To see that our method is still useful even if there is no ground truth, suppose for now that there is no true proportion of documents in each category. Rather, we will suppose that each coder has her own interpretation of the coding scheme, leading to a personal *model* of how documents should be assigned to categories. Under this assumption we can recast our algorithm as a method for interpolating across the coders' models. First, suppose that we are exclusively interested in coder 1's model. Then the evaluation matrix for coder 1 is the identity matrix—by defintion coder 1 has applied her model to the data. The evaluation matrix for coder 2 will have the agreement, a^{12} on the diagonal—where they agree coder 2 is applying coder 1's model—and then the disagreements have to be reassigned to the remaining categories. This will be possible if we can find a matrix with a^{12} down the diagonal $\boldsymbol{E}_{a^{12}}$ such that $\boldsymbol{E}_{a^{12}}^{-1}\bar{\boldsymbol{y}}_2 = \bar{\boldsymbol{y}}_1$. Likewise, if we impose coder 2's model, then the evaluation matrix for coder 2 is the identity model and coder 1's model has the agreement on the diagonal a^{12}.

Now, suppose that we are interested in a mixture between each coder's model of how the documents map into the categories. The coders agree a^{12} of the time, so they share the same model in those instances. Then, for the remaining $(1 - a^{12})$ we will suppose that we impose coder 1's model c_1 share of the documents and coder 2's model $c_2 = (1 - c_1)$ share of the documents. This implies that coder 1's evaluation matrix has $a^{12} + (1 - a^{12})c_1$ on the diagonal and coder 2's evaluation matrix has $a^{12} + (1 - a^{12})(1 - c_1)$. Note, that this implies that the average joint accuracy is $\frac{a^{12}+(1-a^{12})c_1+a^{12}+(1-a^{12})(1-c_1)}{2} = \frac{a^{12}+1}{2}$ or the maximum accuracy in the previous section.

For each c_1 we can find evaluation matrices $E_{c_1}^1$ and $E_{c_1}^2$ such that $\left(E_{c_1}^1\right)^{-1}\bar{\boldsymbol{y}}_1 = \left(E_{c_1}^2\right)^{-1}\bar{\boldsymbol{y}}_2$. For each category k we can then search over c_1 to obtain maximum and minimum values for each category. This, however, yields the equivalent interval as in Equation 3.3: optimizing over different values of c_1 is equivalent to searching over the set $\mathbb{E}$. Therefore, the interval estimator can also be interpreted as a method for characterizing the potential proportions, given coder disagreement.

3.4 Extending the Algorithm to Include Non-Overlapping Coders and Sampling Uncertainty

It is often the case where a pair of coders will double code a subset of data and the remaining documents are coded by a single coder. We can modify our interval estimator to accomodate this additional classification data. Suppose that the coders double code a subset of documents $D = D_1 \cap D_2$. Define $\bar{\boldsymbol{y}}_D^c$ as the proportion for coder c in each category when applied to set D and define $\bar{\boldsymbol{y}}^c$ as the proportion in each category for all D_c coding decisions—including both double and single coded documents. Our constraint that the evaluation matrices, when inverted, return the same proportion can now only be applied to the subset of documents that are coded by both coders. That is $(E^1)^{-1} \bar{\boldsymbol{y}}_D^1 = (E^2)^{-1} \bar{\boldsymbol{y}}_D^2$, but this need not necessarilly hold in general for $\bar{\boldsymbol{y}}^c$ because of differences in the documents included in coder c's sample–even if coder 1 and 2's evaluation matrices are identical.

To include the single-coded documents, we have to introduce an additional assumption: that our coders make the same kind of errors in the double-coded and the single-coded documents. We will call this the "evaluation matrix stability assumption".

Assumption 3. *__Evaluation Matrix Stability Assumption__ Call coder c's evaluation for the double-coded documents $\boldsymbol{E}_D^c$ and for the single-coded documents $\boldsymbol{E}_{D_c}^c$. We will suppose that $\boldsymbol{E}_D^c = \boldsymbol{E}_{D_c}^c = \boldsymbol{E}^c$*

Assumption 3 is quite likely to hold in practice, particularly if the double-coded documents are a random sample from the entire collection of documents. We will define the set $\mathbb{E}$ as in Section 3.2, but only require the constraint to hold over the double-coded documents. Using these assumptions and optimizing over $\mathbb{E}$, Corollary 1 provides an interval estimator that uses all the coded data.

Corollary 1. *Suppose that coder 1 and coder 2 have agreement a^{12} and Assumptions 2, 1, and 3 hold. Define $\bar{\pi}_k^{int}$ as*

$$\bar{\pi}_k^{int} = \left[\min_{\tilde{E}^1, \tilde{E}^2 \in \mathbb{E}} \frac{(\tilde{E}^1)^{-1}\bar{\boldsymbol{y}}^1 + (\tilde{E}^2)^{-1}\bar{\boldsymbol{y}}^2}{2}\Big|_k, \max_{\tilde{E}^1, \tilde{E}^2 \in \mathbb{E}} \frac{(\tilde{E}^1)^{-1}\bar{\boldsymbol{y}}^1 + (\tilde{E}^2)^{-1}\bar{\boldsymbol{y}}^2}{2}\Big|_k \right]$$

Then $\bar{\pi}_k \in \bar{\pi}_k^{int}$

Proof. Under Assumptions 2, 1, and 3 the true evaluation matrices $(E^1, E^2) \in \mathbb{E}$. And if E^1 and E^2 are the true evaluation matrices across both double and single-coded documents, then, $\bar{\pi}_k = \frac{(E^1)^{-1}\bar{\boldsymbol{y}}^1 + (E^2)^{-1}\bar{\boldsymbol{y}}^2}{2}\Big|_k$ ☐

Thus far we have assumed that we have estimates of the population agreement between coders and proportions. Of course, in any sample our estimates of those parameters will also be noisy. To capture the uncertainty from that noise we can perform a simple bootstrapping procedure. Specifically, we perform a vanilla bootstrap at the double-coded document level and then apply our algorithm. We can then obtain conservative bounds on the proportions by taking the minimum of the minimum of each category across the boot strap iterations, and the maximum of the maximum of each category of the boot strap iterations.

No. Coded	Bootstrap	Simulation Type	Proportion Contained
		Table 2: Simulation Evidence	
		Maximum Validity	
100	No	Equal	0.60
100	Yes	Equal	0.93
500	No	Equal	0.93
500	Yes	Equal	1
1000	No	Equal	0.99
1000	Yes	Equal	1
1000	No	Differential	0.96
10000	No	Equal	0.98
10000	No	Differential	0.99
30000	No	Equal	1
30000	No	Differential	0.99
		Independent Coders	
100	No	Equal	0.90
1000	No	Equal	1
10000	No	Equal	1
10000	No	Non-Constant	0.86

3.5   Simulation Evidence for the Bounding Algorithm

To demonstrate the performance of our algorithm we apply it to simulated data, generated under a variety of assumptions about how coders are performing. We then examine the proportion of time our algorithm covers the ground truth of the simulation and how the width of the intervals we construct depends upon the coders' agreement rate.

First consider the top-portion of Table 2. For these simulations we generate our synthetic coding data assuming our coders are performing with maximum joint validity for a varying number of documents (left-hand column). We then report whether we used a bootstrap to estimate the key quantities of interest (second column from left), whether the coders were assumed to have equal validity (or $\epsilon^c = \frac{1+a^{12}}{2}$) for $c = 1, 2$) or with one coder outperforming the other (third column from the left), and finally the proportion of times the estimated interval contains the true proportions. In each of these simulations, we vary the agreement rate from 0.6 to 0.95.

The top-portion of Table 2 shows that under a wide range of settings our interval estimator performs well. When there is a small sample, for example, the uncertainty in estimating the key quantities of interest leads the interval estimator to contain the truth a small number of times, 60% of the time. This, however, is remedied with the bootstrap. As the sample size increases the interval estimator contains the truth almost always. While the Propositions imply that this number should be 1 the interval will sometimes fail to contain the true value because our algorithm is an approximate inference procedure. We can easily improve the approximation by searching over a more granular set of values, at the cost of the algorithm taking longer to complete.

The bottom-portion of Table 2 relaxes the maximum validity assumption, instead supposing that our coders are making independent coding decisions. Note that the assumption of independence is not necessarily a large departure from our assumption of maximum validity. For example, suppose that coders agree on 81% of the documents. If the code independently then we would suppose each coder's validity is 90%, while under maximum joint validity each coder would have validity 90.5%.[1] Perhaps, then, it is not surprising that our algorithm tends to perform well when coders are making independent coding decisions.

The bottom two rows of Table 2 relax the assumption that coders have constant validity across categories. Here, we see that the interval estimator does not perform as well as the other settings. That said, the estimator does often contain the truth, and in the next section we introduce a way to perform a sensitivity analysis that will provide more conservative bounds and therefore improve the coverage of the algorithm.

One might be concerned that the coverage rates in Table 2 are merely the results of creating large intervals that are therefore not particularly informative. Figure 1 shows that this is not the case. In it, we plot the width of all intervals from the top-half of Table 2. The thick-black lines is a local linear regression between the interval width and the agreement rate among coders (where the variability emerges because of the random characteristics of the particular simulation). This demonstrates that for high levels of agreement the intervals are often quite narrow, and yet, still tend to cover the truth regularly.

3.6   Income Inequality News Coverage

Our first application of the algorithm is to an analysis of income inequality news coverage over time. The coded statements come from McCall (2013), which seeks to refute arguments that Americans care little about income inequality. A key component of her analysis is assessment of coverage of income inequality coverage in the news. McCall (2013) employs a team of undergraduate coders to hand code news stories and we examine three categories the coders used: (1) whether the story was irrelevant for the analysis, (2) whether the story was about inequality, and (3) an aggregated category capturing whether the story covered stories broadly about changes in the economy. All together the validation set contained 121 total double-coded documents and the two coders have an agreement rate of 0.88.

Table 3 shows the average of the coders classifications across the three categories and provides the bounds both assuming the population variables are correct and for a bootstrapped values. The first two rows shows that both coders place a very similar proportion of documents in each category, reflecting the high agreement rate between the coders. Further, it shows that a relatively small proportion of the articles—only about 11.5%—are about inequality. The next two rows provide the bounds on the proportion in each category, treating the estimates of agreement and from our coders as population parameters. The bounds are fairly narrow, and show that between 7% and 12% of the articles are about inequality.

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[1]Of course, the difference can be larger, depending on the agreement rate.

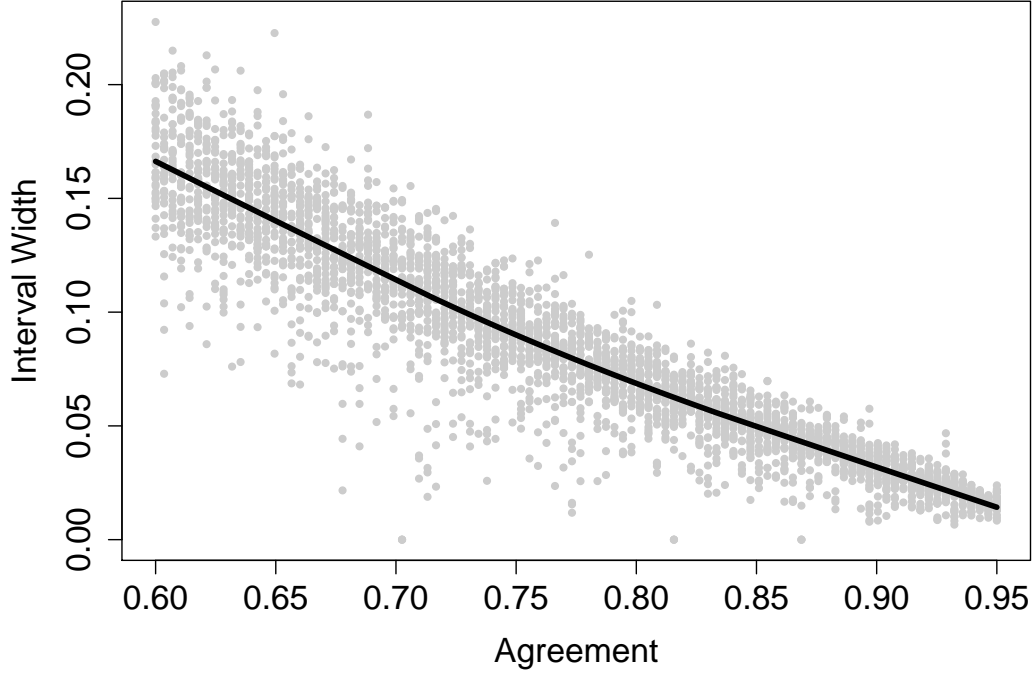Figure 1: Interval Width Declines as Coders Agree More

Table 3: Bounds on the Proportion of News Stories about Inequality

Method	Irrelevant	Inequality	Economy/Changes
Coder 1	0.46	0.12	0.41
Coder 2	0.47	0.11	0.42
Bounds, No Bootstrap, Min	0.46	0.07	0.41
Bounds, No Bootstrap, Max	0.49	0.12	0.44
Bounds, Bootstrap, Min	0.37	0.00	0.31
Bounds, Boostrap, Max	0.60	0.19	0.54

The bottom two rows, however, show the result of bootstrapping our algorithm 100 times and then forming the bounds. Including the estimation uncertainty in our bounding procedure widens the bounds considerably. In spite of the substantial intercoder agreement, the small number of double coded documents implies that there is still substantial uncertainty about the distribution of documents across the categories.

4   Relaxing the Constant Validity Assumption

We derived our interval estimator in the previous section assuming that our coders make errors at the same rate for all categories. This is, of course, a strong assumption. During coding tasks researchers often have a strong intuition that certain categories are easy for coders to accurately classify documents, while other categories are harder for coders to classify. In this section we develop a methodology that generalizes our method for different rates of errors across categories.

To generalize to the non-constant case, we first write all the diagonal entries of each coder's evaluation matrix as a function of the validity rate for the first category, ϵ_{11}^c.[2] Specifically, we will write the validity rate for category k, ϵ_{kk}^1, as

$$\epsilon_{kk}^1 \quad = \quad b_k^1 \epsilon_{11}^1 \tag{4.1}$$

In Equation 4.1 b_k^1 represents the rate of errors in category k relative to the rate in category 1 for coder 1, or the coefficient for category k. If $b_k^1 > 1$ then coder coder 1 is more accurate when classifying documents that belong to category k than category 1 and if $b_k^1 < 1$ then coder 1 is less accurate for category k.

We can collect the coefficients for each coder c into $\boldsymbol{b}^c = (1, b_2^c, \ldots, b_K^c)$. The vector $\boldsymbol{b}^c$ can accommodate any collection of validities across categories, so writing the validities in this way is merely a way to reparameterize the validities without imposing additional assumptions.

While individual coders may have their own idiosyncratic set of errors for each category, we make a simplifying assumption: that each coder has the same set of coefficients. Formally, this supposes that $\boldsymbol{b}^1 = \boldsymbol{b}^2 = \boldsymbol{b}$. Substantively, this supposes that coders' validities are proportional across categories, *not* that the validities are identical across categories. Further, note that this assumption generalizes the constant validity assumption in the previous section, where $\boldsymbol{b} = (1, 1, \ldots, 1)$. Assumption 4 states the assumption formally:

Assumption 4. *Coders 1 and 2 have proportional validities across categories. Formally, we suppose that $\boldsymbol{b}^1 = \boldsymbol{b}^2 = \boldsymbol{b}$.*

Assumption 4, while not strictly necessary, will be extremely helpful in deriving a version of our algorithm with different validity rates for different categories.

In order to apply our algorithm we have to make some assumptions about the values of $\boldsymbol{b}$. Our first assumption will be the most restrictive: that the researcher

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[2]Note, that this is without loss of generality. We are merely using the first category as a benchmark and it implies no other assumptions about the categories.

knows the values of $\boldsymbol{b}$. We will first derive our interval given this assumption, then we will use the derivation from that case to make the weaker assumption that we can rank the values of each b_k, even if we do not know the specific values.

To start, suppose that we know the values of each b_k and we make the maximum average validity assumption. From the proof of Proposition 1 we know that the following equation holds:

$$a^{12} \;=\; \mathrm{P}(\text{Coder 1 Correct}) + \mathrm{P}(\text{Coder 2 Correct}) - 1 \tag{4.2}$$

If we know the values of $\boldsymbol{b}$ then we can rewrite this to obtain:

$$a^{12} + 1 \;=\; \underbrace{\epsilon^1_{11}\bar{\pi}_1 + \epsilon^1_{22}\bar{\pi}_2 + \ldots + \epsilon^1_{KK}\bar{\pi}_k}_{\mathrm{P}(\text{Coder 1 Correct})} + \underbrace{\epsilon^2_{11}\bar{\pi}_1 + \epsilon^2_{22}\bar{\pi}_2 + \ldots + \epsilon^2_{KK}\bar{\pi}_k}_{\mathrm{P}(\text{Coder 2 Correct})}$$

$$= \;\; \epsilon^1_{11}\underbrace{(\bar{\pi}_1 + b_2\bar{\pi}_2 + \ldots + b_K\bar{\pi}_K)}_{g} + \epsilon^2_{11}\underbrace{(\bar{\pi}_1 + b_2\bar{\pi}_2 + \ldots + b_K\bar{\pi}_K)}_{g} \tag{4.3}$$

where g is a constant that is the result of the inner product of $\boldsymbol{b}$ and $\boldsymbol{\pi}$. Simple algebra then reveals that the average value of each coder's validity for category 1 is $\frac{1+a^{12}}{2g}$. Note, that under constant validity $g = 1$, yielding the prior formula for the maximum average validity. Of course, we do not know the values of g because it requires knowing the true values of $\bar{\boldsymbol{\pi}}$. But, we can use the other assumptions to obtain maximum and minimum values for g, which we can subsequently use to infer the potential validity values.

To place bounds on g first note that if maximum average validity holds then when coders agree, they are correct. Call $\tilde{y}_k$ the proportion of the time coder 1 and coder 2 both place a document into category k. The quantity $\tilde{y}_k$ is useful because it represents a lower bound on the proportion of documents in category k. Further the maximum in category, k based solely on coder agreement, is $1 - \sum_{j \neq k} \tilde{y}_j$. Based on these bounds, the biggest potential value of g would occur if the category with the largest $\boldsymbol{b}$ coefficient, $b_{\max}$, also has the largest proportion of documents possible under the constraints. Likewise, the smallest possible value of g occurs if the largest possible proportion of documents, under the constraint, are assigned to the category with the smallest $\boldsymbol{b}$ value, $b_{\min}$. Further, because each validity is between zero and 1 we know that the maximum possible values of $\epsilon^1_{11}, \epsilon^2_{11} = \min\left(\frac{1}{b_{\max}}, 1\right)$.

Together, this information provides a range of values for $g \in [g_{\min}, g_{\max}]$, which imply a parallel set of values of ϵ^1_{11} and ϵ^2_{11} such that $\epsilon^1_{11} + \epsilon^2_{11} = \frac{1+a^{12}}{g}$. Because we know $\boldsymbol{b}$, values of ϵ^1_{11} and ϵ^2_{11} provide the diagonal value for a pair of evaluation matrices. And just like before, we can use the other properties that evaluation matrices must satisfy and collect all such pairs for all potential values of g into $\mathbb{E}_{\boldsymbol{b}}$. We can then define our new interval estimator as,

$$\bar{\pi}^{\mathrm{int},\boldsymbol{b}}_k \;=\; \left[\min_{\tilde{E}^c \in \mathbb{E}_{\boldsymbol{b}}} \left(\tilde{\boldsymbol{E}}^c\right)^{-1} \bar{\boldsymbol{y}}^c|_k, \; \max_{\tilde{E}^c \in \mathbb{E}_{\boldsymbol{b}}} \left(\tilde{\boldsymbol{E}}^c\right)^{-1} \bar{\boldsymbol{y}}^c|_k \right]$$

16

In Proposition 3, we prove that if $\boldsymbol{b}$ is known or if an ordering of the diagonal elements of the evaluation matrices is known, then the bounds will contain the true proportions.

Proposition 3. *Suppose Assumption 1 is true.*

1) *If Assumption 4 is true, then $\bar{\pi}_k \in \bar{\pi}_k^{int,\boldsymbol{b}}$.*

2) *Suppose an ordering of the accuracy rates for each category is known, which implies that an ordering for the components of $\boldsymbol{b}$ is known. Define $\boldsymbol{B}$ as the set of coefficients consistent with that ordering. Then,*

$$\bar{\pi}_k \in \left[\min_{\tilde{\boldsymbol{b}} \in \boldsymbol{B}} \bar{\pi}_k^{int,\tilde{\boldsymbol{b}}}, \max_{\tilde{\boldsymbol{b}} \in \boldsymbol{B}} \bar{\pi}_k^{int,\tilde{\boldsymbol{b}}} \right]$$

Proof. As in the proof of Proposition 2, we show that the assumptions imply the true evaluation matrices are in the set of evaluation matrices considered. To show this for (1) note that the assumptions imply that the true evaluation matrices, $\boldsymbol{E}^1, \boldsymbol{E}^2 \in \mathbb{E}_{\boldsymbol{b}}$. This implies that $\bar{\pi}_k \in \bar{\pi}_k^{int,\boldsymbol{b}}$. Likewise, the assumptions are sufficient if we search over all possible coefficients consistent with the ordering, which is exactly what is contained in $\boldsymbol{B}$. Therefore, $\bar{\pi}_k \in \left[\min_{\tilde{\boldsymbol{b}} \in \boldsymbol{B}} \bar{\pi}_k^{int,\tilde{\boldsymbol{b}}}, \max_{\tilde{\boldsymbol{b}} \in \boldsymbol{B}} \bar{\pi}_k^{int,\tilde{\boldsymbol{b}}} \right]$. □

5 Relaxing the Maximum Joint Accuracy Assumption

Our analysis has relied upon the assumption that our coders are performing as accurately as possible given the level of agreement. Of course this could be a strong assumption. Coders are likely to make the same error or they may disagree and both be incorrect. In this section we describe a sensitivity analysis that relaxes the assumption of maximum average validity, providing a method for examining lower levels of joint validity.

Proposition 1 shows that the maximum average validity for the coders is $\frac{1+a^{12}}{2}$. The minimum average validity is always 0 (assuming $K > 2$ and that we have two coders). To relax the assumption of maximum joint validity, we can fix a level of joint validity $\gamma \in [0, \frac{1+a^{12}}{2}]$ and then examine the potential validity values at that level of joint validity. Proposition 4 shows that at a wide range of values of γ, the agreement rate still provides information about the range of values for coder 1 and 2's validity.

Proposition 4. *Suppose $\dot{\epsilon} = \gamma$, coder 1 and coder 2 have agreement rate a^{12}, $K > 2$, and Assumption 2. Then if $\gamma \in [\frac{1-a^{12}}{2}, \frac{1+a^{12}}{2}]$, $\epsilon^1 \in [\gamma - \frac{1-a^{12}}{2}, \gamma + \frac{1-a^{12}}{2}]$ and $\epsilon^2 = 2\gamma - \epsilon^1$. If $\gamma \in [0, \frac{1-a^{12}}{2}]$ then $\epsilon^1 \in [0, 2\gamma]$ and $\epsilon^2 = 2\gamma - \epsilon^1$.*

Proof. Without loss of generality, consider ϵ^1 and fix $\gamma \in [\frac{1-a^{12}}{2}, \frac{1+a^{12}}{2}]$. Consider first the upper bound. The upper bound on ϵ^1 occurs if coder 1 is always correct

when the coders disagree, which occurs $1 - a^{12}$. If this is true then coder 2 is only correct when she agrees with coder 1. This implies

$$\epsilon^1 = \epsilon^2 + (1 - a^{12})$$
$$2\gamma = \epsilon^1 + \epsilon^2$$

where the second equation follows from the definition of γ. Solving for ϵ^1 yields $\epsilon^1 = \gamma + \frac{1-a^{12}}{2}$. Note that at the maximum $\epsilon^2 = \gamma - \frac{1-a^{12}}{2}$.

While this same argument also provides the lower bound we will now derive it explicitly. Note that ϵ^1 is small as possible if coder 1 is wrong in all instances she disagrees with coder 2. This implies that $\epsilon^2 - (1 - a^{12})$ of the time coder 1 agrees with coder 2 and must be correct. Thus

$$\epsilon^1 = \epsilon^2 - (1 - a^{12})$$
$$2\gamma = \epsilon^1 + \epsilon^2$$

Solving for ϵ^1 yields $\epsilon^1 = \gamma - \frac{1+a^{12}}{2}$.

If $\gamma \in [0, \frac{1-a^{12}}{2})$, then the agreement rate provides no additional constraint on the average validity, because the value of γ is sufficiently small that the coders need not have any overlapping agreement. □

Proposition 4 shows that for a wide range of joint validity, the agreement rate provides information about how much the coders can disagree and therefore specifying a level of joint validity implies a range for the individual coders' validity. Proposition 4 provides a useful sensitivity analysis because it generalizes the upper bound in Proposition 1. To see this, note that the at the maximum joint validity $\gamma = \frac{1+a^{12}}{2}$. The maximum value for ϵ^1 is therefore $\epsilon^1 = \gamma + \frac{1-a^{12}}{2} = \frac{1+a^{12}}{2} + \frac{1-a^{12}}{2} = \frac{2}{2} = 1$. And the lower bound on $\epsilon^1 = \frac{1+a^{12}}{2} - \frac{1-a^{12}}{2} = \frac{2a^{12}}{2} = a^{12}$.

Figure 5 provides a visualization of how the range of joint validity that is possible under different levels of agreement. The horizontal axis in each plot is coder 1's accuracy, ϵ^1 and the vertical axis in each plot is coder 2's accuracy, ϵ^2. The grey area in each plot defines the set of possible values for our coder's accuracy for all the potential values of γ. The thick black lines in each plot define iso-joint validity lines where $\epsilon^1 + \epsilon^2 = 1 + a^{12}$ (line closest to the top-right corner) and $\epsilon^1 + \epsilon^2 = 1 - a^{12}$ (line closest to the origin). We can represent other iso-joint validity lines with the equation $\epsilon^2 = 2\gamma - \epsilon^1$.

Consider the plot in the top-left corner, which shows the range values when the coders disagree. In this case, all values in the lower-half of the unit square are possible. For all potential values of γ under this agreement rate $\gamma = \frac{1+a^{12}}{2} = 0.5$, $\gamma[0, 0.5]$ the range of validity values for both coder 1 and coder 2 spans from 0 to 1. The top-right plot shows that as agreement increases, we have more information about the possible joint-validity values. And as the bottom right-plot shows, that as $a^{12} = 1$, then for any value of γ then only $e_1 = e_2 = \gamma$ is possible.

We use Proposition 4 and Figure 5 to relax the assumption that our coders are performing at maximum average validity. Our previous algorithm proceeds selecting the maximum value of $\gamma = \frac{1+a^{12}}{2}$. Setting lower values of γ then relaxes our optimistic assumption, allowing our coders to agree and be incorrect and for instances where

18

Figure 2: Geometric Interpretation of the Maximum Validity Assumption Suggests
a Sensitivity Analysis
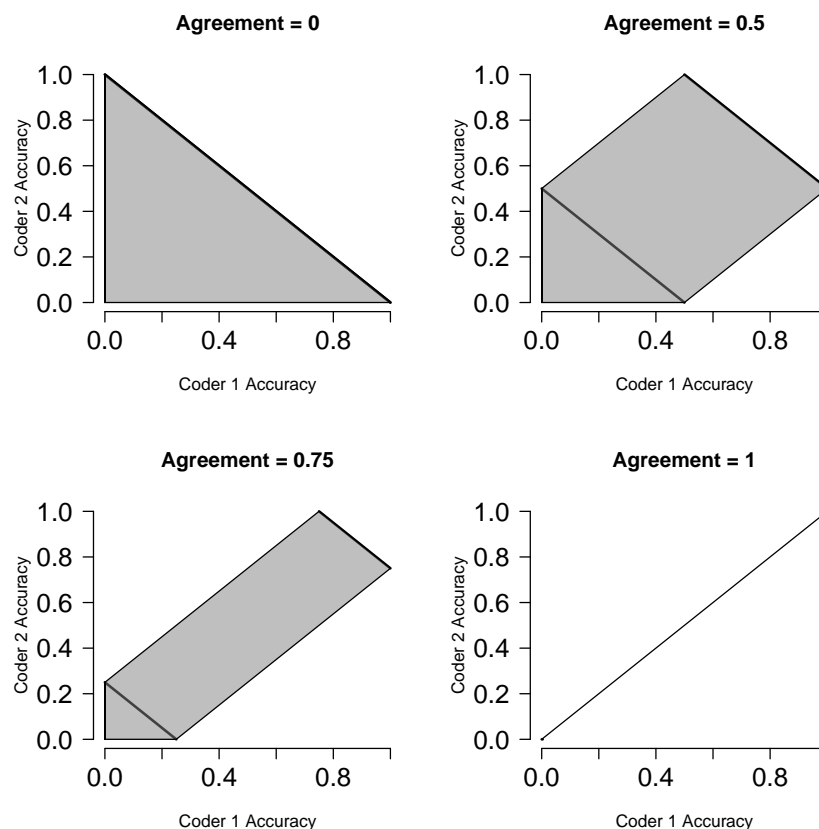
This figure provides a geometric interpretation of the relationship between agreement and accuracy. Note that as the agreement rate is lowered a wider range of accuracies are considered for each coder. This implies that examining how the interval increases as the agreement rate decreases provides a means of assessing the sensitivity of our assumption that our coders are performing as accurately as possible.

they disagree and one coder is not correct. For any level of γ we can obtain a set of potential evaluation matrices with the appropriate joint accuracy, (E_γ^1, E_γ^2). Imposing the constraints, that $\left(E_\gamma^1\right)^{-1} \bar{\boldsymbol{y}}^1 = \left(E_\gamma^2\right)^{-1} \bar{\boldsymbol{y}}^2$ and that $\left(E_\gamma^c\right)^{-1} \bar{\boldsymbol{y}}^c \in \Delta^{K-1}$ for all c. Collect the pairs of matrices that satisfy the constraints for a given level of γ in the set $\mathbb{E}_\gamma$. For a given γ, the interval estimator is then:

$$\bar{\pi}_k{}^\gamma = \left[\min_{\tilde{E}^c \in \mathbb{E}_\gamma} \left(\tilde{E}^c\right)^{-1} \bar{\boldsymbol{y}}^c, \max_{\tilde{E}^c \in \mathbb{E}_\gamma} \left(\tilde{E}^c\right)^{-1} \bar{\boldsymbol{y}}^c \right] \tag{5.1}$$

where the difference between Equation 3.3 and Equation 5.1 the set of matrices considered at different values of γ. If we suppose that the coders average validity $\dot{\epsilon} \in [\gamma_{\text{low}}, \gamma_{\text{high}}]$, then we consider a range of potential γ values. We call this our $\tilde{\gamma}$-level average validity assumption,

Assumption 5. $\tilde{\gamma}$-*level average validity* Coders have average validity $\dot{\epsilon} \in [\gamma_{low}, \gamma_{high}]$ where $\gamma_{high} \leq \frac{1+a^{12}}{2}$ and $\gamma_{low} \geq 0$.

We can encode this broader set of joint validities by optimizing over this interval of gammas, $\pi_k^{\tilde{\gamma}}$,

$$\bar{\pi}_k^{\tilde{\gamma}} = \left[\min_{\gamma \in \tilde{\gamma}} \bar{\pi}_k^\gamma, \max_{\gamma \in \tilde{\gamma}} \bar{\pi}_k^\gamma \right]$$

Proposition 5 shows that $\pi_k^{\tilde{\gamma}}$ will contain the true proportions.

Proposition 5. *Suppose Assumption 2 and 5 hold. Then $\bar{\pi}_k \in \pi_k^{\tilde{\gamma}}$.*

Proof. Assumptions 2 and 5 imply that the true evaluation matrices $(\boldsymbol{E}^1, \boldsymbol{E}^2) \in \mathbb{E}_{\tilde{\gamma}}$ and therefore $\bar{\pi}_k \in \pi_k^{\tilde{\gamma}}$. $\qquad\square$

5.1 Comparing Our Method to State of the Art Parametric Models for Coder Annotation

An alternative approach in Computer Science and Natural Language Processing use parametric models to adjust for potential biases in coders' decisions (Li, Yu and Zhou, 2013; Zhang et al., 2014; Zhou et al., 2014; Passonneau and Carpenter, 2014). While there are many variants of models in the literature, they share a common framework first introduced in Dawid and Skene (1979). Dawid and Skene (1979) model the decisions of annotators using a mixture model, estimated with an Expectation Maximization algorithm.

Like the framework in this paper, Dawid and Skene (1979) suppose that $\bar{\boldsymbol{\pi}}$ represents the true proportion in each category. Given a value of $\bar{\boldsymbol{\pi}}$ the true label for each document is then drawn from a Multinomial distribution $\pi_d \sim \text{Multinomial}(1, \bar{\boldsymbol{\pi}})$. Conditional on the true label for each document, the model then supposes that each coder generates her label with a draw from a Multinomial distribution, with $y_d^c \sim \text{Multinomial}(1, \boldsymbol{\epsilon}^c{}_{\pi_d})$ where $\boldsymbol{\epsilon}^c{}_{\pi_d}$ refers to the π_d^{th} column of the evaluation matrix.

If we assume flat priors on $\bar{\boldsymbol{\pi}}$ and $\boldsymbol{\epsilon}^c_{\pi_d}$ and collect all C evaluation matrices into $\boldsymbol{E}$, we obtain the following posterior distribution:

$$p(\bar{\boldsymbol{\pi}}, \boldsymbol{\pi}, \boldsymbol{E}|\boldsymbol{Y}) \quad \propto \quad \prod_{d=1}^{D}\prod_{k=1}^{K}\left[\bar{\pi}_k \prod_{c=1}^{C}\prod_{j=1}^{K}(\epsilon^c_{jk})^{I(y^c_d=j)}\right]^{I(\pi_d=k)} \tag{5.2}$$

To obtain MAP estimates of Equation 5.2 a straightforward EM algorithm is used (Passonneau and Carpenter, 2014). Other work attempts to introduce assumptions to improve the learning rates of the estimates or the starting values for optimization (Li, Yu and Zhou, 2013; Zhang et al., 2014; Zhou et al., 2014).

Like other mixture models, Equation 5.2 has many local optima and, as a result, it is highly sensitive to the initial starting values used to fit the model. If the model assumptions fit and starting values close to the true proportions are used, then the model can provide accurate estimates of the true proportions. Even in this setting, however, there are no guarantees that the estimates and credible intervals will cover the truth. Further, it remains unclear how best to deal with the values from many different starting values and how to incorporate that uncertainty in quantities of interest.

But in settings where coders' decisions do not fit the assumptions of the model— or where the coders' may make systematic errors—then the Dawid and Skene (1979) model will be less useful. As we now demonstrate, even after we apply many different starting values the model will provide inferences that are systematically biased away from the truth, except in special cases.

To examine the behavior of the Dawid and Skene (1979) model, we simulated 100 different coding tasks. Specifically, we suppose that our coders agree 60% of the time, each has constant validity of 60%, codes 1000 documents, that $\gamma = 0.6$ (implying that each coder is correct 60% of the time) and that the true labels for each document are drawn according to, $\pi_d \sim \text{Multinomial}(1, (0.6, 0.3, 0.1))$. For each individual simulation we then applied the Dawid and Skene (1979) model 100 times. To apply the Dawid and Skene (1979), we set the starting values in two ways: (1) we used the proportion of documents in each category on the subset of documents where the coders agree as starting values for the proportions and random values for the evaluation matrix, which we label the `Coder Agree` procedure and (2) random values for both the proportion in all categories and the evaluation matrices, which we label the `Random Start` procedure. We used the subset of documents the coders agreed on as starting values and allowed the evaluation matrix starting values to be determined randomly.

The results of applying the Dawid and Skene (1979) model for all three categories is shown in Table 4. The first column of Table 4 examines the coverage behavior for the Dawid and Skene (1979) estimator. For each initialization procedure, we use the 100 restarts to define a range of values for each of the proportions. And the first column shows that in only a small percentage of the cases will creating an interval using different starting values from the Dawid and Skene (1979) will cover the true value. This low coverage rate maintains even though the width of the interval obtained using random restarts has similar width to the intervals from

our method. Further the second column shows that the Dawid and Skene (1979) estimator is systematically biased, yielding proportions that deviate systematically from the truth.

Table 4: Comparing Bounds to the Dawid and Skene (1979) Estimator

Model Assumptions	Coverage	Bias			Average Interval Width (Restarts)
DS, Coder Agree	0.16	-0.07	0.06	0.01	0.02
DS, Random Start	0.43	-0.14	0.08	0.06	0.23
Bound, Max Validity	0.68	-	-	-	0.15
Bound, $\gamma \in [0.7, \max]$	0.99	-	-	-	0.23
Bound, $\gamma \in [0.6, \max]$	1.00	-	-	-	0.33

The bottom three rows of Table 4 shows that our interval estimator has superior coverage, even when we make incorrect assumptions about the validity rates. If we assume our coders have maximum validity, then our estimator still covers the true values 68% of the time. If we suppose that our coders have a γ-average validity of 0.7—higher than the true value of 0.6—the interval estimator covers the truth 99% of the time, with an interval width comparable to the width from the Dawid and Skene (1979) model using random restarts. Finally, if we assume the correct level of average joint validity, the interval covers the truth 100% of the time.

Our partial identification approach, then, provides both conceptual clarity about the assumptions necessary to learn from coder disagreement. It also provides superior performance to parametric modls of coder disagreement.

5.2 Examining the Proportion of Stories About Income Inequality

In Section 3.6 we reanalyze double-coded from McCall (2013) and find that sampling variability can result in wide intervals from our method, even when there is a relatively high rate of intercoder agreement. In this section we again analyze McCall (2013) and we show that even if we assume there is no sampling variability, but relax the assumption of maximum validity, there can still remain considerable uncertainty about the true proportion of documents in each category.

Figure 3 shows how the intervals we obtain over the same set of documents from McCall (2013) vary as we consider a wider range of potential levels of joint validity. At the far right of each plot is the interval assuming maximum joint validity, $\frac{1+0.88}{2} = 0.94$ and those intervals replicate the intervals obtain from assuming no sampling variability in our analysis in Section 3.6.

As we move to the left in Figure 3, we consider intervals under lower levels of average joint validity. It is immediately clear that interval width increases substantially as we included instances where our coders are more error prone. And at the far-left of the plot, where we assume the coders are correct 70% of the time average, we obtain the widest intervals for each category. The interval for the Inequality category is [0.42, 0.67]; the interval for the inequality category is [0.0, 0.16]; and the for economy changing category we obtain [0.3, 0.57]. Again, even in a setting where

Figure 3: Relaxing the Assumption of Maximum Joint Validity Yields Wider Intervals
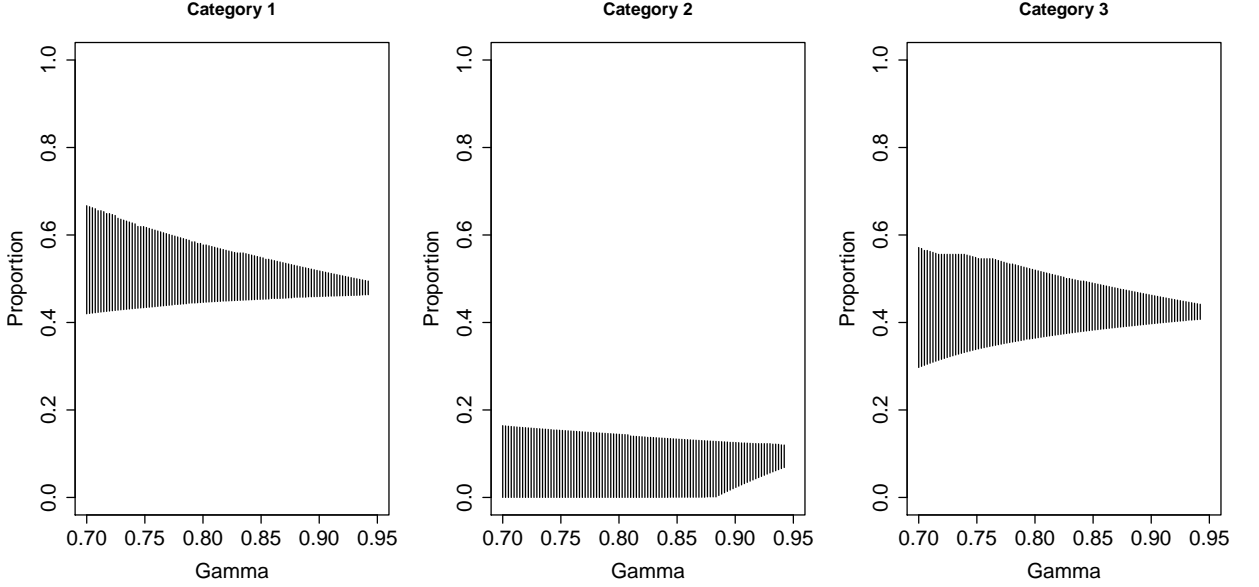
there is a relatively high amount of intercoder reliability, we have certainty about the estimates only if we are willing to assume our coders are performing close to the maximum average joint validity.

6 Uncertainty With More than Two Coders for Each Document

We have so far focused on developing intervals when we have two coders, because this is the most common and often most affordable method used to assess the reliability and validity of document codes. In this section we show how adding additional coders to code a single document can provide even more informative bounds on coders' maximum possible accuracy. This more informative bound arises because our assumption of the maximum possible joint validity and the addition of coders allows us to potentially identify instances where some some coders are *wrong*, allowing us to bound the maximum accuracy of a coder away from 1.

To examine the case with more than two coders, we first introduce some additional notation. For simplicity (and without loss of generality) suppose that our C coders have coded D documents into one-of-K categories. Define the K^C set of potential labels for each document in the set $\mathcal{T}$, with one instance for document d as $\boldsymbol{y}_d \in \mathcal{T}$. The C-element long vector $\boldsymbol{y}_d$ collects the coding decisions across coders and y_d^c describes the coding decision for coder c on document d. We will collect the coding decisions into an $C \times D$ matrix $\boldsymbol{Y}$.

To establish our bounds on coder accuracy with multiple coders, we will examine aggregations of our coder's decisions. An equivalent view of the coding process is that we are asking our coders to vote on the label for a particular document, where we tally the codes from our coders as votes for a document's label. Formally, we define the function $v : \boldsymbol{T} \to \mathbb{C}^K$ as a function that performs this aggregation. For any vector $\boldsymbol{y}_d$ we will say that coder c is in the plurality for document d if $y_d^c = \max_k v(\boldsymbol{y}_d)$ and $|\max_k v(\boldsymbol{y}_d)| = 1$: coder c is in the plurality when her label for document d agrees with the most popular label for document d adn there is a single most-popular label for document d. We will denote the proportion of times that coder c is in the plurality with $p^c = \text{mean}_d \left(I(y_d^c = \max_k v(\boldsymbol{y}_d), |\max_k v(\boldsymbol{y}_d)| = 1) \right.$. We will similarly say that coder c is part of a plurality tie if the number of categories at the maximum is 2 or more, or $|\max_k v(\boldsymbol{y}_m)| > 1$ and $y^c d \in \max_k v(\boldsymbol{y}_d)$. We can similarly define the proportion of times that coder c is part of a plurality tie, $t^c = \text{mean}_d \left(I(y_d^c \in \max_k v(\boldsymbol{y}_d), |\max_k v(\boldsymbol{y}_d)| > 1) \right)$.

Before stating our result, we first generalize our assumption of maximum joint validity for our coders given their agreement. Specifically, we will say assume that our coders have the maximum average validity given the agreement across coders. Formally, we will assume that $\dot{\epsilon} = \text{mean}_c (\epsilon^c)$ is as large as possible, given the coding decisions $\boldsymbol{Y}$. This is a direct generalization from the case where $C = 2$ our Assumption 1, so restate the assumption here:

Assumption 6. *Wisdom of the Coders (General)* *We will suppose that* $\dot{\epsilon} = \text{mean}_c (\epsilon^c)$ *is at a maximum, given* $\boldsymbol{Y}$.

Proposition 6 establishes the bound on coder validity with a larger set of coders.

Proposition 6. *Suppose that there are C coders who code D documents and Assumptions 2 and 6 hold. Then,*

$$\epsilon^c \in [p^c, p^c + t^c]$$

where p^c is the proportion of times c is among the clear plurality of coders and t^c is the proportion of times c is among the pluralities in a plurality tie.

Proof. We first consider the lower bound, given maximum average validity. The assumption that our coders have the maximum average validity given the level of agreement implies that in instances where all the coders agree that the coders are correct. The coders can have two different patterns of disagreement. There can either be a clear plurality of voters for a category, or there will be a plurality tie. Assume first there is a clear plurality of coders. Then our assumption of maximum joint validity implies that those coders are correct. (If not, then average validity could be improved by making them correct without affecting the agreement). This implies that at worst $\epsilon^c = p^c$. To determine the upper bound suppose that coder c is in a plurality tie t^c of the time. Our assumption of maximum average validity and the coder's agreement provides no information about which group in the plurality tie is correct. Therefore, coder c's accuracy is highest if each instance she is involved in a plurality tie she is correct. This implies that maximum value of $\epsilon^c = p^c + t^c$. Thus $\epsilon^c \in [p^c, p^c + t^c]$. $\square$

Proposition 6 shows that with additional coders we can obtain a more informative upper bound on a coder's decision. The extra information arises because our assumption of maximum accuracy given the agreement level among coders implies that any instance where a plurality of voters agree, they must be correct. This implication is very similar to *Wisdom of the Crowds* results. Proposition 6 generalizes the upper bound from Proposition 1. If $C = 2$, then $p^c = a^{12}$ and $t^c = (1 - a^{12})$. If we assume that coder 1 is always correct when they disagree with coder 2 then the upper bound on coder 1's accuracy is $\epsilon^1 = 1$.

While there is more information from coders that could be used to incorporate information into our algorithm, our empirical case studies have shown us that this creates an overly restrictive set of assumptions that often implies an empty interval. Instead, we use the results in Proposition 6 to apply a modified version of our algorithm to each pair of coders. Suppose that we have a collection of coders C coders and that each coder c and coder j's validity lies in the interval from Proposition 6. For each pair of coders c and j we can define the new pairs of evaluation matrices, $\tilde{E}^c, \tilde{E}^j \in \mathbb{E}^{cj}$ such that $\left(\tilde{E}^c\right)^{-1} \bar{y}^c \in \Delta^{K-1}$, $\left(\tilde{E}^j\right)^{-1} \in \Delta^{K-1}$ and $\left(\tilde{E}^c\right)^{-1} \bar{y}^c = \tilde{E}_j^{-1} \bar{y}_j$, and the diagonal elements satisfy the interval from Proposition 6 and $\epsilon^c + \epsilon^j \leq p^c + p^j + t^c + t^j - t^{cj}$ where t^{cj} is the proportion of the time coder c and coder j are in the plurality tie together. We can search over all pairs of coders c, j to define our new interval,[3]

$$\bar{\pi}_k^{\text{int}} = \left[\min_{i,j} \min_{\tilde{E}_i \in \mathbb{E}_{ij}} \tilde{E}_i^{-1} \bar{y}_i|_k, \max_{i,j} \max_{\tilde{E}_i \in \mathbb{E}_{ij}} \tilde{E}_i^{-1} \bar{y}_i|_k\right] \tag{6.1}$$

6.1 Analyzing the Credit Claiming Statements in House Press Releases

Grimmer, Westwood and Messing (2014) analyze the rate of credit claiming across members of the US House. Grimmer, Westwood and Messing (2014) use a collection of 789 triple-coded press releases as a training set for a supervised learning procedure. Here, we analyze only the hand coded press releases to better understand the uncertainty from hand coding. We work with a modified version of their data, analyzing three categories of press releases: advertising, credit claiming for money, and all other press releases. The first three rows in Table 5 presents the proportion of press releases in each of the categories from each of the coders. The relatively close alignment across the categories reflects the high rate of agreement among the coders. All three coders agreed 68.1% of the press releases; coder 1 and 2, while coder 3 disagreed in 9.3% of the press releases; coder 1 and coder 3 agreed while coder 2 disagreed in 7.2% of the press releases; coder 2 and coder 3 agreed, while coder 1 disagreed in 13.7% of the press releases; and all three coders disagreed in only 1.6% of the press releases. We use this agreement rate as input to obtain bounds on the proportion in each category, following Equation 6.1.

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[3]Equation 6.1 will provide a conservative estimate of coder uncertainty, because it fails to include all the information from the multiply coded data that might further restrict the range of validity that we search over.

Table 5: Reanalyzing the Rate of Credit Claiming in a Sample of US House Press Releases

Coder Number	Advertising	Credit Claiming	Other
1	0.29	0.21	0.51
2	0.23	0.25	0.51
3	0.32	0.30	0.38
Bounds			
Minimum	0.20	0.20	0.51
Maximum	0.26	0.24	0.56

7 Propagating Uncertainty from Coder Disagreement to Other Parameter Estimates

The proportion of documents that lie in a set of categories is often an intermediate outcome of interest. The true quantity of interest is often the effect of some intervention on the prevalence of that document or how the prevalence of some category of interest affects some other outcome of interest. In this section we describe a straightforward method for incorporating our uncertainty as a result of our coders' disagreement into the next stage of an analysis. For simplicity we will focus on a linear regression with a regression coefficient as the quantity of interest, but our method extends to other parametric models and more elaborate quantities of interest in a straightforward way. Proposition 8 in the Appendix shows this straightforward generalization.

Suppose that our team of coders classifies documents from S sources with the true proportion in category k from source s given by $\bar{\pi}_k{}^p$. A source might be a politician, a survey respondent, a country's leaders, a year a document was written, or any other covariate stratum. Rather than observe a K component vector $\bar{y}^c$ for each coder c, we now obtain a $K \times S$ matrix of proportions for each coder, $\bar{Y}^c$. We denote the sth row of this matrix for the cth coder as $\bar{y}_s^c$. This row provides the proportion of observations in each category for a particular source of interest.

We will suppose that our interest is in understanding the relationship between a vector of covariates for each source X_s—which we collect into X—and its relationship with the prevalence of category k across the S sources. We will suppose the following data generating process:

$$\bar{\pi}_k{}^s = \beta X_s + \nu_s$$

where ν_p is independent and identically distributed errors, with $E[\nu_s|X_s] = 0$. Our quantity of interest is β or the conditional relationship between covariates prevalence of category k across sources. If we observe $\bar{\pi}_k{}^s$ for each source s then the usual regression estimator will be an unbiased and consistent estimator of β.

As we have argued throughout this paper, it is unlikely that our coders have performed without error. The consequences of coder error for our regression can be substantial. If our coders code with error, then we can write the observed vector of proportions for category k $\bar{y}_k$ as $\bar{y}_k = \pi_k + \delta_k$, where δ_k is the vector of average

bias across coders[4] from coding for each source. Unless $\delta_k^s = 0$ for each source s the regression estimator will no longer be consistent or unbiased because:

$$\left(\boldsymbol{X}'\boldsymbol{X}\right)^{-1}\boldsymbol{X}'\bar{\boldsymbol{y}} = \left(\boldsymbol{X}'\boldsymbol{X}\right)^{-1}\boldsymbol{X}'\bar{\boldsymbol{\pi}} + \left(\boldsymbol{X}'\boldsymbol{X}\right)^{-1}\boldsymbol{X}'\boldsymbol{\delta}$$

$$= \boldsymbol{\beta} + \left(\boldsymbol{X}'\boldsymbol{X}\right)^{-1}\boldsymbol{X}'\boldsymbol{\delta} \tag{7.1}$$

The second term remains so long as our coders code with error. While the regression will be biased and inconsistent, we can develop an interval estimator that will contain the true value of the regression coefficient of interest. To do, we first have to make an additional assumption about the errors our coders make.

When focused on the proportion of documents in each category, it is true by construction that there is one matrix that relates our coders' decisions to the true proportions. When we examine many sources, however, it is possible that coders make different sorts of errors from different sources. We will denote the errors that coder c makes on documents from source s as $\boldsymbol{E}_s^c$. The specific errors that coder c makes on source s might be the result of coders' biases toward the source—such as liberal college students coding documents from Republicans—or the result of some sources using more ambiguous language—which might occur with language from historical sources.

In many instances, however, coders are likely to make the same errors across sources. This might be particularly likely if coders are randomly assigned to documents that are from a relatively similar set of sources—say official documents from members of Congress—and unnecessary information is removed to limit the possibility that coders' biases will affect their decisions.

To develop our interval estimator for the coders' proportions, we will make the additional assumption that our coders make the same errors across sources. Specifically we will assume that $\boldsymbol{E}_s^c = \boldsymbol{E}^c$ for all sources s.

Assumption 7. *Source Evaluation Matrix Stability Assumption* *For all sources s, we assume that $\boldsymbol{E}_s^c = \boldsymbol{E}^c$ for all coders c.*

In other words, our coders can make their own specific errors, based on their confusion about the coding rules or their attention to the documents, but across sources, s, the coders make the same errors—they aren't modifying their errors according to the source of the document they are coding. This assumption, in addition to likely being true, will substantially simplify the development of an interval estimator for quantities of interest. Under this assumption, then, implies that for the true evaluation matrix $\boldsymbol{E}^c$, $\left(\boldsymbol{E}^{-1}\right)^c \bar{\boldsymbol{y}}_s^c = \bar{\boldsymbol{\pi}}^s$ for all sources s.

To construct the interval estimator, suppose our quantity of interest is the tth coefficient β_t. We will assume again that our two coders have the maximum joint validity given the level of agreement. We will further restrict the set of evaluation

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[4]This use of the term bias is different than what can be found sometimes in the literature on handcoding. Di Eugenio and Glass (2004) for instance use this term to indicate the degree to which the coders disagree. Another example from Byrt, Bishop and Carlin (1993): If Observers A and B differ in their assessment of the frequency of occurrence of a condition in a study group, we say that there is a bias between the observers. (page 424).

matrices to be just those that, when inverted and applied to the estimated proportions $(\boldsymbol{E}^c)^{-1}\bar{\boldsymbol{y}}^c$ the result lies in the simplex and where $(\boldsymbol{E}^1)^{-1}\bar{\boldsymbol{y}}^1 = (\boldsymbol{E}^2)^{-1}\bar{\boldsymbol{y}}^2$. Define $\mathbb{E}$ as the set of pair matrices that satisfy these conditions. Then we can define an interval estimator for β_t^{int} as,

$$\beta_t^{\text{int}} = \left[\min_{(\boldsymbol{E}^1) \in \mathbb{E}} \left(\boldsymbol{X}'\boldsymbol{X} \right)^{-1} \boldsymbol{X}' \left((\boldsymbol{E}^1)^{-1}\bar{\boldsymbol{y}}^1 \right) |_t, \max_{(\boldsymbol{E}^1) \in \mathbb{E}} \left(\boldsymbol{X}'\boldsymbol{X} \right)^{-1} \boldsymbol{X}' \left((\boldsymbol{E}^1)^{-1}\bar{\boldsymbol{y}}^1 \right) |_t \right]$$

Where, again $|_t$ selects the tth element from the vector.

Proposition 7 shows that, β_t will be contained in β_t^{int}.

Proposition 7. *Suppose that Assumptions 2, 1, and 7 hold for all categories and all coders. Suppose that using the true proportions $\bar{\boldsymbol{\pi}}$ and covariates $\boldsymbol{X}$ yields the regression coefficient is β_t. Then $\beta_t \in \beta_t^{int}$.*

Proof. If coders have maximum average validity, then the true evaluation matrices $(\boldsymbol{E}^1, \boldsymbol{E}^2) \in \mathbb{E}$. This implies that $\left(\boldsymbol{X}'\boldsymbol{X} \right)^{-1} \boldsymbol{X}' (\boldsymbol{E}^1)^{-1}\bar{\boldsymbol{y}}^1 = \boldsymbol{\beta}$ and $\left(\boldsymbol{X}'\boldsymbol{X} \right)^{-1} \boldsymbol{X}' (\boldsymbol{E}^2)^{-1}\bar{\boldsymbol{y}}^2 = \boldsymbol{\beta}$. Therefore $\beta_t \in \beta_t^{\text{int}}$ $\square$

In Appendix C we show that our interval estimator generalizes for any estimator that would be consistent if the true proportions were used as either a dependent or independent variables.

7.1   Analyzing the Taunting Rate Over Time

Grimmer, King, and Superti (2015) (GKS) analyze the rate US Senators engage in taunting: explicit, public, and negative attacks on the other party or its members. To measure the rate of taunting GKS engages in a massive hand coding effort, employing a team of coders to label thousands of Senate speeches. GKS ensured that 10% of all the speeches were double coded, to measure the agreement of the coders. In this section we use a subset of the hand-coded data to examine how the rate of taunting has varied over time and to demonstrate the need to adjust for coder disagreement.

To apply the algorithm from the previous section, we aggregate their more granular coding into three categories: taunting, policy discussion, and other types of Senate speech. We then focused on the proportion in each category from the two most prolific coders in the data set. This yielded 379 double coded Senate speeches and the two coders had an overall agreement rate across the three categories of 0.83. Altogether the two coders classified and 11,136 total speeches coded.[5] Using the complete set of codes we measured the proportion of speeches from each senator in each of three categories for the two coders.

Using the agreement rate between the coders and the measures of the proportion in each category, we used the procedure in the previous section to measure the average taunting rate from the 101st to the 109th Congress. The thick-black line in

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[5]The total number of double-coded speeches is substantially lower than 10% because there were many coders included in the data set and here we focus on only one pair of coders.

Figure 4 shows the intervals on the average taunting rate in each Congress from our bounding procedure, while the dashed line demonstrate the average taunting rate if we fail to propogate coder uncertainty.

Figure 4: Taunting Rates in the US Senate, Across Congresses
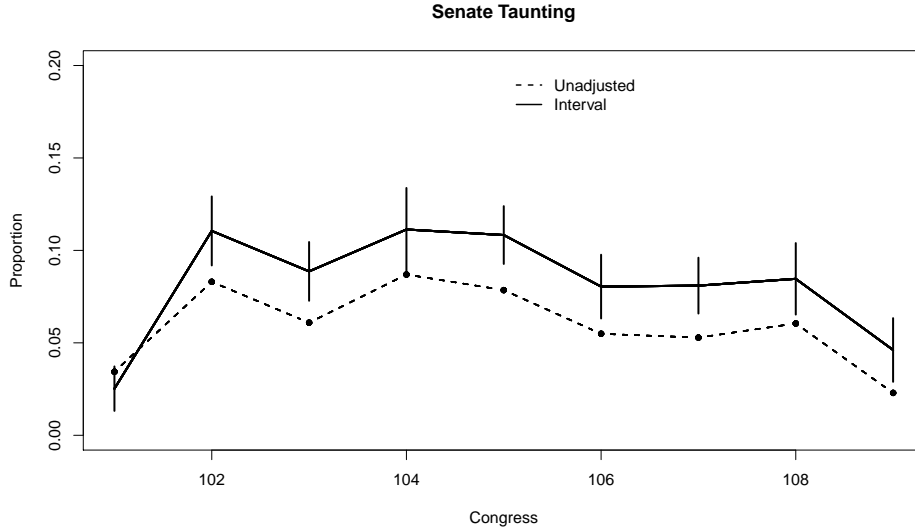
Senate Taunting

Figure 4 demonstrates that failing to include coder error can lead to a serious underestimation of the taunting rate in the Congress. In the worst cases, the unadjusted taunting rate is more than a full percentage point lower in the *minimum* of the interval from the bounding procedure. Overall, the unadjusted taunting rate underestimates the taunting rate about 0.5 percentage points—or 9 percent underestimate.

Substantively, Figure 4 shows that there has not been a secular increase in taunting in the Senate. Rather, taunting appears highest immediately after the Republican revolution, but subsided somewhat during the Bush administration.

8 Conclusion

In this paper we have addressed a fundamental issue in the analysis of hand coded data: how to ensure coder disagreement is incorporated in our analyses. We introduce a set of behavioral assumptions about how coders perform their task that yield interval estimators that are guaranteed to cover the true values if the assumptions hold. Applying our method to standard hand-coding exercises reveals that there is substantial uncertainty about the true value of proportions, even when coders have seemingly high levels of agreement.

Beyond the applications discussed here, our procedure has a wide range of applications that we have not yet considered. For example, it can be useful for providing a non-parametric method for aggregating insights from ensemble classification methods. Our focus on coder error is also useful for competitive learning tasks, where machine classification methods are compared to a human "gold-standard" coding to determine a state of the art coding method.

There are also several important issues in applying and using our method that deserve further attention. Developing a more efficient optimization procedure will make our approach easier to apply. Further, extending the procedure to a large number of categories will require additional assumptions on the evaluation-matrix, but information that is likely available in the coder decisions.

A   Summary of Notation

- Define $\text{mean}_j(a_j) = \sum_{j=1}^{J} a_j / J$ where $J = |\{j : \forall j\}|$.

- indices:

  d $(d = 1, \ldots, D)$ document

  s $(s = 1, \ldots, S)$ source (of documents)

  k $(k = 1, \ldots, K)$ category

  c $(c = 1, \ldots, C)$ coder (always as superscript)

- truth:

  $\pi_d \in \{1, \ldots, K\}$ true category for document d (always as subscript)

  $\bar{\pi}_k = \text{mean}_d[I(\pi_d = k)]$ true proportion of documents in category k, with vector $\bar{\boldsymbol{\pi}} = \{\bar{\pi}_k : k = 1, \ldots, K\}$

- measures:

  $y_d^c \in \{1, \ldots, K\}$ coder c's classification of document d

  $\bar{y}_k^c = \text{mean}_d[I(y_d^c = k)]$, proportion of documents classified into category k by coder c, the average over all coders $\bar{y}_k = \text{mean}_c(\bar{y}_k^c)$, and vector $\bar{\boldsymbol{y}} = \{\bar{y}_k : k = 1, \ldots, K\}$

- Validity and Reliability:

  Evaluation Proportion: ϵ_{jk}^c are misclassification proportions, the proportion of times coder c classifies a document into category j among those whose true category is k, with matrix $\boldsymbol{E}^c = \{\epsilon_{jk}^c\}$, the diagonals of which, ϵ_{jj}^c, are the **validities**

  Confusion Matrix element: $m_{jk}^{12} = \text{mean}_d[I(y_d^1 = j, y_d^2 = k)]$, with diagonal elements being the **reliabilities (or agreement proportions)**, $a^{12} = \sum_{k=1}^{K} m_{kk}^{12}$

- Concepts

  Prior, Standard Assumptions: (a) above some (mysterious) threshold of reliability, validity is not effected (i.e., the estimator is not biased); (b) at least one coder is always right

  Constant Category Validity Assumption: $\epsilon_{kk}^c = \epsilon^c$ for all k

Wisdom of the Coders: Technically, this is a "Maximum Average Coder Validity" assumption, or $\dot{\epsilon}^{12} = \frac{1+a^{12}}{2}$. Equivalently, as we show, this means if a plurality of coders of document d agree, they vote for the truth; if there is a tie, then at least one side votes with the truth.

Coding Generation Process: $\bar{y}_k^c = \sum_{k=1}^{K} \epsilon_{jk}^c \pi_k$, or in matrix form: $\bar{y}^c = E^c \pi$

Coder Trust Assumption: $\epsilon^1 = \frac{1+a^{12}}{2} + \lambda \frac{1-a^{12}}{2}$, $\epsilon^2 = 1 + a^{12} - \epsilon^1$ for $\lambda \in [-1, 1]$

B   Algorithm To Obtain Identification Region

To obtain the intervals using coding decisions and our assumptions, we use a brute force approach to optimization. Specifically, suppose that we observe decisions from coder 1 $\boldsymbol{y}^1$ and coding decisions from coder 2 $\boldsymbol{y}^2$, with agreement rate a^{12}. From Proposition 1 that $\epsilon^1 + \epsilon^2 = 1 + a^{12}$.

Using this information we perform a grid search over the agreement values and the evaluation matrices, while encoding constraints. Our algorithm first searches over the possible accuracy levels, given agreement. For each accuracy level we then perform a grid search over the evaluation matrices consistent with that level of accuracy. By iterating over combinations of accuracy and evaluation matrices we are able to characterize the maximum and minimum possible values.

More specifically, we use the following procedure. Suppose we have a K category measure.

- For each ϵ^1 and ϵ^2 such that $\epsilon^1 + \epsilon^2 = 1 + \mathtt{a}^{12}$:

    - For each column k and each $\boldsymbol{z}_k \in \Delta^{K-1}$:

        - Create matrix E^2 such that:

$$
\begin{pmatrix}
\epsilon^2 & (1-\epsilon^2)z_{12} & \cdots & (1-\epsilon^2)z_{1K} \\
(1-\epsilon^2)z_{21} & \epsilon^2 & \cdots & (1-\epsilon^2)z_{2K} \\
\vdots & \vdots & \ddots & \vdots \\
(1-\epsilon^2)z_{K1} & (1-\epsilon^2)z_{K2} & \cdots & \epsilon^2
\end{pmatrix}
\tag{B.1}
$$

        - For each column k and each $\boldsymbol{x}_k \in \Delta^{K-2}$ create matrix E^1 such that

$$
\begin{pmatrix}
\epsilon^1 & (1-\epsilon^1-\tilde{x}_2)x_{12} & \cdots & (1-\epsilon^1-\tilde{x}^K)x_{1K} \\
\tilde{x}_1 & \epsilon^1 & \cdots & \tilde{x}_K \\
(1-\epsilon^1-\tilde{x}_1)x_{31}) & \tilde{x}_2 & \cdots & (1-\epsilon^1-\tilde{x}^K)x_{3K}) \\
\vdots & \vdots & \ddots & \vdots \\
(1-\epsilon^1-\tilde{x}_1)x_{K1} & (1-\epsilon^1)x_{K2} & \cdots & \epsilon^1
\end{pmatrix}
\tag{B.2}
$$

        where each $\tilde{x}_k$ are unknown values, leaving us with K unknown values, which we collect into $\tilde{\boldsymbol{x}}$

- Solve for $\tilde{x}_k$ such that

$$(\boldsymbol{E}^1)^{-1}\bar{\boldsymbol{y}}^1 - (\boldsymbol{E}^2)^{-1}\bar{\boldsymbol{y}}^2 \;\; = \;\; \boldsymbol{0} \tag{B.3}$$

using a Newton-Raphson algorithm. This ensures that the proportions are equal.
- If there is a solution use it to calculate $\boldsymbol{E}^1$ and then if $(\boldsymbol{E}^1)^{-1}\bar{\boldsymbol{y}}^1 \in \Delta^{K-1}$ then update the following for each k

$$\begin{aligned}
\min_{\mathbf{k}} &= \min(\min_{\mathbf{k}}, [(\boldsymbol{E}^1)^{-1}\bar{\boldsymbol{y}}^1]_k) \\
\max_{\mathbf{k}} &= \max(\max_{\mathbf{k}}, [(\boldsymbol{E}^1)^{-1}\bar{\boldsymbol{y}}^1]_k)
\end{aligned}$$

- For each k return $\min_{\mathbf{k}}$ and $\max_{\mathbf{k}}$

Modifying the algorithm for the cases described above is straightforward. To perform the sensitivity analysis we alter the range of validity that we search over. When one coder is trusted more, we restrict the set of matrices that we search over. When we have multiple coders we restrict the range of joint validity that we search over.

The algorithm becomes increasingly complex as the number of categories increase. But for three and four categories the model is easily fit and extended for all of our cases. In many examples one or two categories are of interest and therefore more complex schemes could be reduced to a 2-3 category scheme.

C   A General Approach to Using Proportions as Dependent and Independent Variables

In Section 7 we demonstrated that our approach could be applied to a case where a parameter from a linear regression is the quantity of interest. In this section we provide a straightforward extension of Proposition 7 to any consistent estimator where the estimated proportions may be the dependent or independent variable.

For this general case, suppose that the set of potential covariate values for the sources is $\boldsymbol{\mathcal{X}}$. Suppose we have an estimator $g : \Delta^{K-1} \times \boldsymbol{\mathcal{X}} \to \Re$, $g(\boldsymbol{\pi}, \boldsymbol{X})$ for some quantity of interest ϕ. Note that we are being intentionally ambiguous about whether $\boldsymbol{\pi}$ is the independent or dependent variable. We will suppose that $g(\boldsymbol{\pi}, \boldsymbol{X})$ is a consistent estimator of ϕ, so that $g(\boldsymbol{\pi}, \boldsymbol{aX}) \to^{\text{plim}} \phi$. As in Section 7, we also restrict attention to only those evaluation matrices that, when inverted and applied to the hand coded estimate, returns estimates in the simplex and where the inverted evaluation matrices, applied to the coders' estimates, provide the value. As before, define this set of pairs of matrices as $\mathbb{E}$.

Then, we can define an interval estimator for the quantity of interest ϕ^{int} as,

$$\phi^{\text{int}} \;\; = \;\; \left[\min_{(\boldsymbol{E}^1, \boldsymbol{E}^2) \in \mathbb{E}} g((\boldsymbol{E}^1)^{-1} \bar{\boldsymbol{y}}^1, \boldsymbol{X}), \max_{(\boldsymbol{E}^1, \boldsymbol{E}^2) \in \mathbb{E}} g((\boldsymbol{E}^1)^{-1} \bar{\boldsymbol{y}}^1, \boldsymbol{X}) \right] \tag{C.1}$$

Proposition 8 shows that Equation C.1 will contain the true value of ϕ.

Proposition 8. *Suppose that $g(\boldsymbol{\pi}, \boldsymbol{X})$ is a consistent estimator for ϕ, that coder 1 and coder 2 have maximum joint validity, that the coders' errors are source independent, and that the coders have constant validity across categories. Then $\phi \in \phi^{int}$.*

Proof. The argument parallels the proof of Proposition 8. First note that the true evaluation matrices $\boldsymbol{E}^1, \boldsymbol{E}^2 \in \mathbb{E}$ by Assumptions 2, 1, and 3. This implies that $\left(\boldsymbol{E}^1\right)^{-1} \bar{\boldsymbol{y}}^c = \left(\boldsymbol{E}^2\right)^{-1} \bar{\boldsymbol{y}}^c = \boldsymbol{\pi}$ and thus $g(\left(\boldsymbol{E}^1\right)^{-1} \bar{\boldsymbol{y}}^c, \boldsymbol{X}) = \phi$ and $g(\left(\boldsymbol{E}^2\right)^{-1} \bar{\boldsymbol{y}}^2, \boldsymbol{X}) = \phi$. Therefore $\phi \in \phi^{\text{int}}$ $\qquad\qquad\square$

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