

Finding Convincing Views to Endorse a Claim

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ABSTRACT

Recent studies investigated the challenge of assessing the strength of a given claim extracted from a dataset, particularly the claim's potential of being misleading and cherry-picked. We focus on claims that compare answers to an aggregate query posed on a view that selects tuples. The strength of a claim amounts to the question of how likely it is that the view is carefully chosen to support the claim, whereas less careful choices would lead to contradictory claims. We embark on the study of the reverse task that offers a complementary angle in the critical assessment of data-based claims: given a claim, find useful supporting views. The goal of this task is twofold. On the one hand, we aim to assist users in finding significant evidence of phenomena of interest. On the other hand, we wish to provide them with machinery to criticize or counter given claims by extracting evidence of opposing statements.

To be effective, the supporting sub-population should be significant and defined by a "natural" view. We discuss several measures of naturalness and propose ways of extracting the best views under each measure (and combinations thereof). The main challenge is the computational cost, as naïve search is infeasible. We devise anytime algorithms that deploy two main steps: (1) a preliminary construction of a ranked list of attribute combinations that are assessed using fast-to-compute features, *and* (2) an efficient search for the actual views based on each attribute combination. We present a thorough experimental study that shows the effectiveness of our algorithms in terms of quality and execution cost. We also present a user study to assess the usefulness of the naturalness measures.

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The source code, data, and/or other artifacts have been made available at https://github.com/shunita/claimendorse.

1 INTRODUCTION

Data collected rigorously can be used to identify important phenomena in an arguably objective manner. Hence, evidence from data is often used to qualify stated claims. Yet, for this reason, data-based arguments involve considerable risks. Cherrypicking, for instance, refers to basing the claim on a query with specific components that may seem natural, but are nevertheless crucial for supporting the claim rather than its opposite. The risk also goes in the other direction: basing a claim on the general population may give the false impression that the claim holds in important subpopulations (an extreme manifestation of this falsehood is the Simpson's Paradox). Past work proposed ways of assessing whether a given claim, commonly phrased as an aggregate query over a database, is cherrypicked [3, 4, 33, 34]. This can be seen as a branch in the more general area of computational fact-checking [15, 17, 20, 26, 62, 63, 68]. Relevant techniques involve machine learning [33], query perturbation [3, 4, 62, 63], and Natural Language Processing (NLP) [15, 20].

In this work, we study this task in the reverse direction: given a claim that is false in the database, we aim to find natural views where the claim holds. We refer to this problem as claim endorsement. The objective is to enrich people's machinery with a complementary tool for the critical assessment of claims. In particular, effective claim endorsement can help users better understand the mechanism of cherry-picking and its involved risks. It can also allow users to look for queries that weaken or invalidate a stated claim; for example, a user can look for alternative phrasings that seem just as natural, yet support the opposite claim on the same database. Such an exercise can further help users assess how amenable a particular dataset is to support contradictory claims, and how suspicious we should be, in general, regarding allegations drawn from this dataset. Such abilities may be useful in many domains. For example, they may be used to closely inspect the claims of political candidates, or explore the extent of social issues. They could also be applied in journalism, academic research, and various fields where critical analysis of data-driven claims is crucial.

EXAMPLE 1.1. Consider the Stack Overflow Developers Survey dataset¹ that includes information about Hi-Tech workers such as backgrounds, demographics, and annual salaries. In this dataset, the average salary of people with a master's degree is *lower* than that of people with only a bachelor's degree. Alex, a social scientist, aims to challenge this observation. She may do so to understand the potential of cherry-picking by a malicious party, to explore the extent of the phenomenon, or to find out whether the dataset can still support the master's degree. Hence, she seeks support

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¹https://survey.stackoverflow.co/2022 (accessed Oct 2024)

for the claim that *individuals with a master's degree earn more, on average, than those with a bachelor's degree.* She finds that the claim holds for people in the field of Data Science and Machine Learning (Subpopulation I). It also holds for people in Germany. In addition, she finds less compelling subpopulations, like people who use Zoom for office communication (Subpopulation II), and people who do not know the size of their organization.

The example illustrates that, to be effective, the views that endorse the claim should be as significant and "natural" to support for the claim. As the notion of "natural" subpopulations is subjective, we incorporate various measures of naturalness. These measures are drawn from different fields such as information theory, NLP, and statistics. For instance, drawing from past proposals [53, 60], one simple measure evaluates the subpopulation's size, with a larger size indicating broader applicability of the claim. Another measure assesses the claim's strength within a subpopulation, determined through tests of statistical significance [58]. Additionally, naturalness can be inferred based on the linguistic relationship between the target attribute and the predicate that defines the view, using language models such as sentence transformers [45] (as we adopt in our implementation). Our proposed framework is designed to identify and extract the highest-scoring views under each measure, as well as combinations of measures, for a given user claim.

EXAMPLE 1.2. We continue Example 1.1, and focus on Subpopulations I and II. The definition of Subpopulation II is, arguably, more artificial and arbitrary, hence less convincing; Subpopulation I appears more natural and supportive of the claim. This intuition is captured by various measures of naturalness. First, Subpopulation I is larger (1446 individuals vs. 743). Second, the criterion of "office synchronous communication tool" is quite unexpected due to lack of connection to salaries and/or academic degrees; we aim to capture this difference through measures from statistics and NLP. Third, within Subpopulation I, master's people earn considerably more than bachelor's (average of \$183K vs. \$144K), whereas the advantage in Subpopulation II is modest (\$117K compared to \$112K). These considerations are indeed reflected by lower scores for Subpopulation II according to the naturalness measures that we consider in this paper. See Section 5.2 for additional examples.

In more formal terms, we define (in Section 2) the problem of *claim endorsement* as follows. We are given an SQL query Q with grouping and aggregation (e.g., Average) over a database (relation) D, along with a claim stating that the aggregate value of one group is higher than that of another group. We seek useful *refinements* that restrict Q by adding predicates p to the selection condition of Q, so that the claim holds true in the refined query. Each refinement defines a sub-population where the claim holds, and we seek sub-populations that are *natural* in order to best convince of the claim's validity. Hence, the goal is to compute the top-k refinements according to a function v that quantifies the naturalness of each refinement (in the context of Q and D). As aforesaid, we discuss a collection of basic measures of naturalness that cover different aspects of what "natural" can be. Hence, a system can solve multiple instances of the problem with different measures.

The main technical challenge that we address (Section 4) is the high computational cost of claim endorsement: refinements can be made out of many attributes, attribute combinations, and value assignments; moreover, each candidate refinement may require costly computation to verify its correctness and measure its naturalness. Efficiency is critical when claim endorsement is used within data exploration, where the user may interactively react to the results of some claims to formulate new ones; response times of days or even hours render the tool ineffective. In particular, listing all possible refinements before ranking is too costly. Similar challenges have been encountered in the search for drill-down and roll-up operators to find the most interesting data parts for exploration [1, 23, 50, 65], the discovery of intriguing data visualizations and explanations [10, 56], and the explanation of outliers in aggregate queries [31, 60]. However, the algorithms in that body of work cannot be applied in our setting where the requirement is to satisfy the claim and, at the same time, achieve high scores of naturalness measures, as they satisfy neither requirement (see Section 6).

Instead of materializing all possible candidates, we devise a framework for *anytime algorithms* that target the incremental generation of high-quality refinements from the very beginning. We instantiate the framework on the aforementioned measures of naturalness. More technically, the algorithms enumerate refinements in a ranked fashion, where the ranking function is, intuitively, well correlated with the naturalness measure, yet efficient to handle. Our framework deploys two main steps: (1) produce a ranked list of attribute combinations according to an easy-to-compute scoring function. (2) for each attribute combination C (in ranked order), compute all value assignments for the corresponding refinements. For each step, we develop several solutions and optimizations that we evaluate in the experiments.

We present a thorough experimental study (Section 5) on three datasets: American Community Survey, Stack Overflow, and a flight-delays dataset. These datasets differ in the number of attributes (from tens to hundreds) and number of tuples (from tens of thousands to millions). We show that our anytime algorithms typically achieve 95% of the quality of the true top-k refinements faster than existing baselines by one-to-two orders of magnitudes. We also present case studies on the three datasets, similarly to Example 1.1.

Additionally, we report a user study that we have conducted to assess how well our proposed naturalness measures align with intuitive concepts, and to compare them to existing solutions. Our study indicates that these measures effectively capture aspects of intuitive naturalness and, furthermore, that they align better with intuition than measures adapted from previous work (e.g., [48]).

In summary, our contributions are as follows: (a) We introduce the claim-endorsement problem. (b) We introduced the concept of naturalness measures for claim endorsement, and proposed several concrete measures, adjusted to the context of this problem. (c) We devise an anytime framework for claim endorsement. (d) We instantiate the framework with algorithms for the proposed measures of naturalness. (e) We present an experimental study that shows the effectiveness of our solutions and analyzes several case studies, and a user study to assess the effectiveness of the naturalness measures.

2 FORMAL FRAMEWORK

In this section, we describe our formal model and define the problem of claim endorsement. We begin with preliminary definitions.

Table 1: Records from ACS [11] used as a running example.

ID	Sex	Occupation	EducationLevel	QoB	Income (K)	
1	F	CS&Math	Bachelor's degree	1	72	
2	F	CS&Math	Master's degree	3	95	
3	F	Education	Master's degree	2	43	
4	F	Sales	High School Diploma	1	35	
5	F	Sales	Bachelor's degree	4	100	
6	М	CS&Math	Bachelor's degree	4	80	
7	М	CS&Math	Master's degree	3	90	
8	М	Education	Master's degree	2	62	
9	М	Sales	Bachelor's degree	1	70	
10	М	Sales	High School Diploma	3	65	

2.1 Databases and Queries

We consider an input database *D* that consists of a single relation with the relation name *R* and the attribute set $Att(D)=\{A_1, \ldots, A_q\}$. Note that this relation can be the result of a query that joins multiple source relations that are outside of the model (as is indeed the case in our experiments with the Flights dataset; see Section 5). We denote by |D| the number of tuples of *D*. By $D[A_{i_1}, \ldots, A_{i_\ell}]$ we denote the (set-semantics) projection of *D* to the attributes $A_{i_1}, \ldots, A_{i_\ell}$. By a slight abuse of notation, for a single attribute A_i we may view $D[A_i]$ as the set of *values* rather than single-value *tuples*.

We assume an aggregate SQL query Q of the following form.

SELECT A_{gb} , $\alpha(A_{agg})$ FROM D WHERE ϕ GROUP BY A_{gb} (1) where $A_{gb} \in \text{Att}(D)$ is the group-by attribute, $A_{agg} \in \text{Att}(D)$ is the aggregate attribute, and α is an aggregate function among Count, Sum, Average, Median, Min and Max. We will refer to a query of this form as a *group-aggregate* query. The result of Q on D, denoted

Q(D), is a set of tuples of the form (g, v), where $g \in D[A_{gb}]$ and

 $v = \alpha(\{t.A_{agg} \mid t \in D \land \phi(t) \land t.A_{gb} = g\}).$

EXAMPLE 2.1. Consider the sample of the ACS dataset [11] shown in Table 1. An analyst may be interested in justifying a Master's degree, so they will first issue the query Q:

SELECT EducationLevel, Average(Income)	
FROM D	
GROUP BY EducationLevel;	

In Q(D), we find that the average income for people with a Master's degree (\$72.5K) is lower than that of Bachelor's degree (\$80.5K).

2.2 Claims and Refinements

Similarly to previous work in the context of query result explanations [31, 35, 47, 53], we study the case where the analyst restricts attention to the relationship between two groups of interest, g_1 and g_2 , in the result Q(D). For these, the analyst may be interested in endorsing a specific claim. We define it formally as follows:

DEFINITION 2.2 (CLAIM). Consider two tuples (g_1, v_1) and (g_2, v_2) in Q(D). We refer to the tuple $\kappa = (g_1, g_2, >)$ as a claim. A groupaggregate query Q' endorses κ (on D), denoted $Q'(D) \models \kappa$, if there are numbers v'_1 and v'_2 such that $(g_1, v'_1) \in Q'(D)$ and $(g_2, v'_2) \in$ Q'(D) and $v'_1 > v'_2$.

We consider the situation where Q violates $(g_1, g_2, >)$, and seek a *refinement* Q' that satisfies it. We focus on refinements that add predicates to the WHERE clause (as done previously, e.g., [33]). EXAMPLE 2.3. Following Example 2.1, the analyst wishes to compare the average income for different degree holders, with the initial assumption that a higher degree implies a higher salary. Yet, she finds that the average income for people with a Master's degree (\$72.5K) is actually *lower* than that of people with a Bachelor's degree (\$80.5K). In our formalism, the analyst is interested in the relationship between $g_1 =$ Master's and $g_2 =$ Bachelor's and their corresponding values in Q(D), namely $v_1 =$ 72.5 and $v_2 =$ 80.5. Hence, the claim (Master's, Bachelor's, >) is violated by Q.

In this paper, we study the problem of searching for query refinements. To that end, we assume a space \mathcal{P} of predicates that can be used to refine the query Q. We consider equality predicates of the form A=v, where $v\in D[A]$, and conjunctions of up to m such predicates, where m is a parameter. This practice has been commonly used in prior query answer explanations research [14, 33, 40, 47, 60]. More formally, we are given a set of *split attributes* from Att(D) that does not include A_{gb} and A_{agg} , and we denote this set by SAtt(D).² Then \mathcal{P} consists of all predicates of the form $A_1 = v_1 \wedge \cdots \wedge A_{\ell} = v_{\ell}$ such that $\ell \leq m$, $A_i \in SAtt(D)$ and $v_i \in D[A_i]$ for all $i = 1, \ldots, \ell$. An *atom* or atomic predicate is a predicate of the form $A_i = v_i$. Given a predicate p of this form, denote by $Att(p) = (A_1, \ldots, A_{\ell})$ the set of attributes used to define p, sorted lexicographically.

DEFINITION 2.4 (REFINEMENT). Let Q be a group-aggregate query as in Equation (1), and let $p \in \mathcal{P}$ be a predicate. The refinement of Qby p is the query Q_p where ϕ is replaced by $\phi' = \phi \wedge p$.

Note that, by Definition 2.2, a refinement Q_p endorses the claim κ if $Q_p \models \kappa$, that is, p selects a subset of D that satisfies $\alpha(g_1) > \alpha(g_2)$.

EXAMPLE 2.5. Recall that in Example 2.3, the query Q violates the claim (Master's, Bachelor's, >). Consider the predicate p_1 given by the expression Occupation = CS&Math. In contrast to Q, the refinement Q_{p_1} satisfies the claim: the average income for Master's degree holders is \$92.5K, yet only \$76K for Bachelor's. Another possible refinement is defined by the predicate p_2 =(QoB=3) (where QoB stands for Quarter of Birth), where v_1 =92.5 and v_2 =65.

Note that our refinements are restricted to conjunctions of equality predicates, in line with past work on explanations that deem such predicates intuitive and understandable [14, 46]. We leave for future work the consideration of a richer class of refinements, including inequality and disjunction, as our space of restricted refinements is already large and computationally challenging.

2.3 Naturalness

Supporting the analyst claim can be performed by finding a certain refinement where the claim holds. However, this refinement should be *natural* in the sense that it should not be overly specific and restricted. For example, the following refinement from the ACS dataset: "people who depart for work between 11:50 and 13:30 and have served in the US military after Sept. 2001" is, arguably, overly specific and can hardly serve as significant support for the claim.

DEFINITION 2.6 (NATURALNESS MEASURE). Let Q be a groupaggregate query, and κ a claim. A naturalness measure (for Q and

²This is similar to prior work on cherry-picking detection [33] and other work on counterfactual explanations, where a subset of attributes cannot be used in the explanation as it is non-actionable [16, 27, 28].

 κ) is a function v that maps pairs (Q_p, D) , where Q_p is a refinement and D is a database, to a numerical score $v(Q_p, D)$.

Semantically, $v(Q_p, D) > v(Q_{p'}, D)$ means that we consider Q_p as more natural than $Q_{p'}$ for the database D (and specifically the results $Q_p(D)$ and $Q_{p'}(D)$). We are not aware of any relevant formalization of naturalness. Intuitively, v aims to quantify (or be well correlated with) the likelihood of a critical listener accepting the claim if it is presented with this refinement.

An example of a naturalness measure is the coverage of Q_p : the fraction of database tuples covered by $\phi \wedge p$. We provide additional examples of measures of naturalness in Section 3 and implement them as part of our framework and empirical study.

2.4 **Problem Definition**

Naturalness is a subjective notion that is unlikely to be expressed precisely by a mathematical formula. In the remainder of the paper, we propose and study several examples of measures that capture what we deem as intuitive aspects of naturalness. Additionally, to broaden the scope of candidates, we consider an output of k refinements instead of a single one, as users are more likely to find the desirable refinements this way even if the measure v does not completely capture their idea of naturalness. Furthermore, good refinements may be found in the *union* of the top-k refinements of *multiple* measures. Hence, we define our main problem as follows.

DEFINITION 2.7 (CLAIM ENDORSEMENT). Fix a predicate space \mathcal{P} and a naturalness measure v. Claim endorsement is the following problem: given a database D, a group-aggregate query Q, a claim $\kappa = (g_1, g_2, >)$ and a natural number k, find k refinements Q_p of Q that endorse κ and have the highest v scores.

This problem definition allows for defining an instance of the problem for each naturalness measure; often, the practical thing to do is to combine several different measures, that is, to retrieve the top-k refinements according to each naturalness measure and to present all of them to the user to select from. Furthermore, the system can define a naturalness measure that combines the measures (e.g., via weighted sum). In our implementation, we also allow for a hyper-parameter M that restricts the refinements to be such that each subpopulation (g_1 and g_2) is of size at least M, so that we avoid refinements that apply to tiny groups. In Section 4, we devise methods to prioritize the search by a combination of all naturalness measures, and we evaluate these methods empirically in Section 5.

EXAMPLE 2.8. Reconsider the dataset in Table 1 and the query computing the average income for each education level, as shown in Example 2.1. An example of an instance of claim endorsement aims to find refinements of up to m=2 atoms so that the aggregated value for g_1 = Master's is higher than the aggregated value for g_2 = Bachelor's (as opposed to the trend on the entire data shown in Example 2.3), and to retrieve the top-20 (k = 20) out of those, according to its coverage as the natural measure of choice. Two of the refinements are described in Example 2.5.

3 EXAMPLES OF NATURALNESS MEASURES

We now present a collection of naturalness measures, adapted from prior work. The examples aim to cover varying intuitive aspects of true naturalness, and are sourced from various domains, including information theory, statistics, and natural language processing. A user can define custom naturalness measures, by implementing $v(Q_p, D)$ and possibly a method to prioritize the search for faster discovery of refinements with high v values. We further discuss custom measures in Section 4.2.

Coverage. This measure is the fraction of tuples covered by the refinement. Recall the condition $\varphi' = \varphi \wedge p$ from the definition of refinement (Definition 2.4). Let $\sigma_{\varphi}(D)$ be the set of tuples from *D* that satisfy φ . We then define $Coverage(Q_p, D) := |\sigma_{\varphi \wedge p}(D)|/|D|$.

Embedding similarity. Modern word embeddings well capture the semantics of textual data [7]. The cosine similarity between the embedding of the predicate (treated as text) and that of the target attribute can measure their perceived relatedness. To compute this measure, a pre-defined mapping $T : A \rightarrow \Sigma^*$ from attribute identifiers to a textual representation is needed. This mapping can be manually defined or extracted from database documentation; in our implementation, we mapped each abbreviation to the full wording. For example, in the ACS dataset, we mapped "MARHM" to "Married in the past 12 months," based on the ACS documentation. An embedding model $E : \Sigma^* \rightarrow \mathbb{R}^d$ emits a *d*-dimensional vector of real numbers for a given textual input. The measure is given by:

$$EmbSim(Q_p, D) \coloneqq CosSim\left(E\left(\bigoplus_{(A_i=v_i)\in p} T(A_i)\oplus T(v_i)\right), E(T(A_{agg}))\right)$$

Here, \oplus represents string concatenation and *CosSim* is the cosine similarity between two vectors. This score aims to measure how natural it is to limit the records according to the predicate $A_i = v_i$, when asking a query about the target attribute. In our implementation, we used a general-purpose sentence-BERT³ model [45].

EXAMPLE 3.1. For p_1 from Example 2.5 we have:

 $EmbSim(Q_{Occupation="CS&Math"}, D) =$

CosSim(E("Occupation CS & Math"), E("Income"))=0.19

For the predicate p_2 , the value of $EmbSim(Q_{QoB=3}, D)$ is

CosSim(E("Quarter Of Birth 3"), E("Income"))=0.07.

This is to be expected since a person's occupation is considered more related to their income than the quarter of the year of birth.

Statistical significance. For this measure, we adopt *hypothesis testing*, which determines whether the difference between group values is significant and indicative of an actual phenomenon. The database is viewed as a sample of a real-world population.

This score has been previously used as a measure of interestingness in the context of insights in views [52]. This measure is defined only when α is Average (where we use the measure $StatSig_{avg}$) or α = Median (where we use $StatSig_{med}$).⁴ When α = Average, the appropriate test is the *two-sided independent T-test* [25]. To define the measure precisely, we denote:

$$\alpha(g_i) \coloneqq \alpha(\{t.A_{agg} | t \in D \land t.A_{gb} = g_i\})$$

Then, the test statistic is given by

 $t^* \coloneqq \frac{\operatorname{Average}(g_1) - \operatorname{Average}(g_2)}{\sqrt{(\operatorname{StdDev}(g_1)^2/\operatorname{Count}(g_1) + (\operatorname{StdDev}(g_2)^2/\operatorname{Count}(g_2))}}$

³Specifically, all-MiniLM-L6-v2: https://huggingface.co/sentence-transformers/all-MiniLM-L6-v2 (accessed Oct 2024)

⁴For the other aggregations, there is no closed-form statistical test. It is possible to use bootstrapping or permutation tests, which we leave for future work.

Under the null hypothesis, the test statistic is t-distributed with df degrees of freedom, where the formula for df is as follows. Let $n_i = \text{Count}(q_i)$ and $s_i = \text{StdDev}(q_i)$. Then,

$$df := \frac{(s_1^2/n_1 + s_2^2/n_2)^2}{(s_1^2/n_1)^2/(n_1 - 1) + (s_2^2/n_2)^2/(n_2 - 1)}$$

The p-value for the two-sided T-test is the probability of observing t^* or a more extreme value, under the null hypothesis (that $X \sim T(df)$). It is given by pvalue := $2 \cdot P(X \ge t^*|X \sim T_{(df)})$.

The naturalness score is defined as the complement of the pvalue. We use the complement instead of the raw p-value so that higher scores will be associated with stronger claims.

$$StatSig_{avg}(Q_p, D) \coloneqq 1 - pvalue$$

For the median difference, we use the median test [51] with Yates correction for continuity [64]. Denote $a_i = |\{t|t.A_{gb} = g_i \wedge t.A_{agg} > Median(g_i)\}|$ and $b_i = Count(g_i)-a_i$. Also, let $EA_i = (a_1+a_2) \cdot (a_i+b_i)/(a_1+b_1+a_2+b_2)$ and $EB_i = (b_1+b_2) \cdot (a_i+b_i)/(a_1+b_1+a_2+b_2)$. The test statistic is given by:

$$x^* := (|a_1 - EA_1| - 0.5)^2 / EA_1 + (|b_1 - EB_1| - 0.5)^2 / EB_1 + + (|a_2 - EA_2| - 0.5)^2 / EA_2 + (|b_2 - EB_2| - 0.5)^2 / EB_2$$
(2)

Under the null hypothesis, the test statistic comes from a Chi-square distribution with 1 degree of freedom. Then the naturalness score is defined as the complement of the probability of observing x^* or a more extreme value, under the null hypothesis (that $X \sim \chi^2_{(1)}$).

$$StatSig_{med}(Q_p, D) \coloneqq 1 - P\left(x \ge x^* | x \sim \chi^2_{(1)}\right)$$

Mutual information (MI). Arguably, predicates that are deemed natural would be based on attributes with some correlation to the target attribute. A possible method to quantify this dependence is MI between attributes, which has been previously used to recommend views for anomaly detection [24]. Given a refinement Q_p , we use MI between the attribute list $Att(p) = (A_1, \ldots, A_\ell)$ that defines p and the target attribute A_{agg} . The definition of MI uses probability distributions over the values of a list of attributes, computed based on their normalized frequencies in the database, and it is given by:

$$MI(Q_p, D) := I((A_1, ..., A_l), A_{agg})$$

= $\Delta_{KL}(P_{((A_1, ..., A_l), A_{agg})} || P_{((A_1, ..., A_l) \otimes P_T)})$

where Δ_{KL} is the KL-divergence distance between probabilities, $P_{((A_1,...,A_l),A_{agg})}$ is the joint distribution of $A_1, ..., A_l$ and A_{agg} , and $P_{((A_1,...,A_l)\otimes P_{Aagg})}$ is their product distribution.

Mutual information is non-negative and unbounded. To maintain the value between [0,1] we normalize all values by dividing them by the largest MI value of an attribute or attribute combination.

Analysis of variance (ANOVA). Another method to quantify the dependence between an attribute combination (A_1, \ldots, A_ℓ) and the aggregate attribute A_{agg} is ANOVA. Our use is similar to what was done by Wu et al. [61] to assess the connection between a view dimension and an outcome dimension.

As in the statistical significance score, the database is viewed again as a sample from a real world population. Given a predicate p, consider the attribute combination used in it: (A_1, \ldots, A_ℓ) . This attribute combination induces a partition of A_{agg} values into θ groups (bags) $\{x_i^1\}_{i=1}^{s_1}, \ldots, \{x_i^\theta\}_{i=1}^{s_\theta}$, where each group is associated

Input : Dataset D, query Q, user claim κ , natural numbers m and
k, a naturalness measure v .
Output : Set <i>M</i> of refinements of <i>O</i> .

1 *Combs* \leftarrow all combinations from SAtt(*D*) of size $\leq m$

2 CombsSorted \leftarrow Sort(Combs, k) // Section 4.2

4 $P \leftarrow \text{Predicates}(\textit{comb}, Q, A_{agg}, A_{gb}, \kappa) // \text{Section 4.1}$

- 5 $M \leftarrow \text{NaturalnessMeasures}(P)$
- 6 Print(M)

with a value combination $(v_1, \ldots, v_\ell) \in D[A_1, \ldots, A_\ell]$:

$$\{x_i^j\}_{i=1}^{s_j} = \{t.A_{agg} | t \in D \land \bigwedge_{p=1}^{t} t.A_p = v_p\}$$

Each group represents a sample from a normally distributed population. The null hypothesis in *ANOVA* is that each of the θ normal distributions has the same mean. The test statistic is based on the averages of each group of numbers in the partition. Let *N* denote the total sample size: $N := \sum_{j=1}^{\theta} s_j$. Let μ denote the total average of A_{aqq} , and denote by μ_j the average of number group *j*.

$$ANOVA(Q_p, D) \coloneqq \frac{\frac{1}{\theta - 1} \sum_{j=1}^{\theta} s_j \cdot (\mu_j - \mu)^2}{\frac{1}{N - \theta} \sum_{j=1}^{\theta} \sum_{i=1}^{s_i} \left(x_i^j - \mu_j \right)^2}$$

The *ANOVA* statistic is non-negative but unbound from above; as with *MI*, we normalize it by dividing it by the largest value in *D*.

Filtering by generality. In OREO [34], the authors propose a method to retrieve counterarguments of a claim and suggest filtering ones contained in others, keeping only the most general. We adapt this concept as follows. Let Q_{p_1} and Q_{p_2} be two refinements comprised of conjunctions of atomic predicates A = v. We say that Q_{p_1} is more general than Q_{p_2} if the set of atoms comprising p_1 is a subset of the set of atoms comprising p_2 . This filter can be composed over any measure of naturalness v, proposed in this section or otherwise. After retrieving the set of refinements and ranking them by v, we can filter only the most general refinements, and present the top-k out of those. We evaluate the effectiveness of this approach through the user study presented in Section 5.3.

4 COMPUTING REFINEMENTS

In this section, we present an *anytime* algorithmic framework to search for the top-scoring refinements. Recall that an *anytime algorithm* can be asked for results at any moment, the quality of the results improves over time, and the goal is to reach close-to-optimal quality early in the computation [69]. Algorithm 1 shows the general approach. We first generate all possible attribute combinations up to a maximal number of *m* atoms (Line 1). Next, we rank the combinations by some prioritization, based on pre-computation and heuristics (Line 2). We then go over the ranked attribute combinations (Line 3) and retrieve the refinements of each combination where the claim holds (Line 4). For each of these, we compute all naturalness measures (Line 5) and print them (Line 6). In the remainder of this section, we describe the functions used in Algorithm 1.

4.1 Finding Predicates for Given Attributes

We first describe Line 4 in Algorithm 1: generating all refinements of a specific attribute combination. Given an attribute combination $(A_1, ...A_l)$, we wish to retrieve all predicates p to endorse the claim, that is, $p = \bigwedge_{i=1}^{l} (A_i = v_i)$ such that $Q_p(D) \models \kappa$. A naive way is to run the following query, that we refer to as *predicate level*, since it is run for each combination of attribute values $v_1, ..., v_l$:

 $\overline{\text{SELECT } A_{qb}, \alpha(A_{agg}) \text{ FROM } D}$

WHERE A_{gb} IN (g_1, g_2) AND $A_1 = v_1$ AND ... AND $A_l = v_l$ GROUP BY A_{ab}

We then verify that $Q_p(D) \models \kappa$, that is, $\alpha(g_1) > \alpha(g_2)$. Nevertheless, that would require the execution of a large number of queries one for each $B \in P(\text{SAtt}(D))$ up to size *m*, for each combination of values of in the database. Instead, we execute one query for each attribute combination. The query returns all value combinations that define refinements that endorse the claim.

The query depends on the choice of aggregate function; we provide an example for the query for α = Average:

```
SELECT A_1, ..., A_\ell
```

 $\begin{array}{l} {\rm AVG}({\rm CASE \ WHEN \ } A_{gb} = g_1 \ {\rm THEN \ } A_{agg} \ {\rm END}) \ {\rm as \ m1}, \\ {\rm AVG}({\rm CASE \ WHEN \ } A_{gb} = g_2 \ {\rm THEN \ } A_{agg} \ {\rm END}) \ {\rm as \ m2}, \\ {\rm FROM \ D} \\ {\rm GROUP \ BY \ } A_1, ..., A_\ell \\ {\rm HAVING \ } {\rm AVG}({\rm CASE \ WHEN \ } A_{gb} = g_1 \ {\rm THEN \ } A_{agg} \ {\rm END}) < \\ {\rm AVG}({\rm CASE \ WHEN \ } A_{gb} = g_2 \ {\rm THEN \ } A_{agg} \ {\rm END}) < \\ {\rm AVG}({\rm CASE \ WHEN \ } A_{gb} = g_1 \ {\rm THEN \ } A_{agg} \ {\rm END}) > M \\ {\rm AND \ COUNT}({\rm CASE \ WHEN \ } A_{gb} = g_2 \ {\rm THEN \ } A_{agg} \ {\rm END}) > M \\ {\rm AND \ COUNT}({\rm CASE \ WHEN \ } A_{gb} = g_2 \ {\rm THEN \ } A_{agg} \ {\rm END}) > M \end{array}$

For each group, we define the two subgroups g_1 and g_2 using the CASE WHEN statements. We found that the use of CASE WHEN yields faster executions than a simple join between a view for each population (an experimental evaluation is included in the full version [2]). The HAVING statement verifies that $\alpha(g_1) > \alpha(g_2)$. We also incorporate (in HAVING) the user-defined threshold on group sizes so that the refinements define sets of size at least M.

On Stack Overflow with m = 1 and Average, predicate-level search takes 10x longer than the attribute-level search (48.5s vs. 4.5s). Predicate-level search uses 842 queries versus only 47 for attribute-level search. See additional comparisons in the full version [2].

As an optimization, we apply pruning based on the group sizes. We pre-compute the maximal group size in a group-by query according to each attribute combination. If that size is at most *M*, we remove the attribute combination from the generated set, since the query above returns an empty result due to the violation of the HAVING statement. Additionally, we preprocess the dataset to count the distinct values of each attribute. We prune attributes with a single distinct value from the generated combinations, since a query refinement using this attribute will have no effect, as all tuples have the same value in this attribute. Lastly, we extend the query above to incorporate the computation of some of the naturalness measures, particularly *Coverage* and *StatSig*. This is done by adding aggregates to the SELECT clause: standard deviation, count, and counts of values above/below the median.

4.2 Prioritization of Attribute Combinations

Next, we propose ways to prioritize attribute combinations, which is done in Line 2 of Algorithm 1.

4.2.1 Prioritization per measure. We differentiate between two types of measures. Attribute-level measures are naturalness measures that are based only on the connection between the attributes $A_1, ..., A_l$ defining the refinement and the aggregate attribute A_{agg} . In our case, these are *MI* and *ANOVA*. For these, the full computation can be done in advance, before knowing the exact query *Q*, the claim κ , or the groups g_1 and g_2 . The computation may be costly, but it is done offline before the analyst issues the query, and can be stored and read when needed. In contrast, the *predicate-level measures* are naturalness measures that are based on the specific predicates that define the refinement, and possibly on the results $\alpha(g_1)$ and $\alpha(g_2)$. Therefore, they cannot be precomputed accurately. In this work, these are the measures *Coverage*, *StatSig*, and *EmbSim*. We will nevertheless propose heuristic methods to estimate these measures based on attribute-level information.

<u>ANOVA and MI</u>. For these, the prioritization is straightforward: we calculate in advance the exact values for each attribute combination, and order by decreasing value.

<u>*EmbSim.*</u> Here, we prioritize combinations by a simplified version of the one defined in Section 3. Define $EmbSimSimple(Q_p, D)$ to be

$$CosSim\left(E\left(\oplus_{(A_i=v_i)\in p} T(A_i)\right), E(T(A_{agg}))\right).$$

This version compares only the attributes that define the predicate (without the values) against the representations of the aggregate attributes. Intuitively, the first string is a part of the string defining the predicate, so we expect *EmbSimSimple* to be close to *EmbSim*.

<u>Coverage</u>. This one prioritizes by (the reverse of) the number of values with over K occurrences for some K (100 in our experiments). We expect attribute combinations that define many large groups to yield large groups where the claim holds.

<u>StatSig</u>. To find predicates with a high *StatSig* score, we wish to quickly identify the attribute combinations that cause the most polarization between the two groups (i.e., high $\alpha(g_1) - \alpha(g_2)$). As a heuristic precomputation, we train two linear-regression models, M_1 and M_2 , on g_1 and g_2 , respectively, to predict the (numeric) value of the attribute A_{agg} . For the split attributes SAtt $(D) = \{A_1, ..., A_q\}$, each model has the form $M(A_1, ..., A_q) = \sum_{i=1}^{q} w_i A_i + c$. The weight w_i is indicative of A_i 's contribution to predicting A_{agg} . Hence, for each A_i , we define $Reg(A_i) = w_i^1 - w_i^2$ where w_i^j is the weight for feature *i* in regression model M_j . We rank combinations by decreasing $Reg(A_1, ..., A_l) = \sum_{i=1}^{l} Reg(A_i)$.

4.2.2 Prioritization for combining all measures. In many cases, a single naturalness measure does not capture all the intuitively natural refinements. Furthermore, the analyst may not know in advance which naturalness measure is best for her query. For these reasons, we next describe prioritization methods that combine single-measure rankings into a single prioritization strategy. For each naturalness measure, we either compute it (or the appropriate heuristic estimate) in advance, or use a fast computation as a preprocessing step, as detailed in Section 4.2.1. We rank the list of attribute combinations according to each computed score. Then, we combine the rankings using one of the following three variations.

<u>Serial</u>: For each individual measure, solve claim endorsement independently and stop when k refinements have been found. If the deadline of Algorithm 1 has not passed, the search continues over the remaining attribute combinations, in their original order.

Merged: Merge the ranked lists in an interleaving fashion. More precisely, we take the first element from each ranking, then the second element from each ranking, and so on. Then we perform the search according to the single merged ranking.

Sampling. A general prioritization heuristic can be done by sampling a portion of the database and finding all refinements that endorse the claim via brute-force search. Note that the time for executing each group-by query on the sample is faster than the full database, so this search is generally faster. (It may also be lossy, since some combinations may be missing.) For each refinement, we compute all measures, relying on either the precomputation for the attribute-level measures (*ANOVA* and *MI*) or the samplebased results for the predicate-level measures. As a heuristic, we define the priority of a combination as the maximum of the average naturalness, over all refinements of the combination.

The advantage of sampling is its independence from the choice of a specific measure. For example, for a custom naturalness measure, if there is a prioritization method for the new measure (like the regression weight for *StatSig*, or the simplified embedding similarity for *EmbSim*), it can be combined with the other rankings. But even without a prioritization method, sampling can still be used for a custom measure, since the process of sampling and searching the sample for predicates for each attribute combination (Section 4.1) is independent of the choice of measures.

5 EXPERIMENTAL EVALUATION

We present experiments to evaluate our framework. We aim to address two main research questions. (Q_1) What is the effectiveness of our framework for claim endorsement? Specifically, what insights are revealed by our framework, and how do they differ from prior work intended for similar (yet different) goals? (Q_2) To what extent do the optimizations contribute to the performance?

Summary. Before we delve into the details of the experiments, we briefly summarize our findings. We observed that our framework generates intuitive and understandable refinements that endorse the claim across various settings. Our approach can generate more convincing refinements than prior work [34, 48]. We found MERGED TOP-k to be the best-performing prioritization. Over all naturalness measures and datasets, MERGED TOP-k was on average 13.4 (and up to 78.3) times faster than the random-order baseline, and on average 16.2 (up to 82.5) times faster than HypDB (as alternative prioritization). Furthermore, MERGED TOP-k method is almost indifferent to the number of tuples and attributes in the database, and to the number of top results (k) and maximal number of atoms (m).

5.1 Experimental Setup

Our code is in Python and publicly available online.⁵ We used the Pandas library for accessing raw dataset files and SQLAlchemy for accessing a PostgreSQL database. We also used sentence transformers [45] ⁶ for the word-embedding model. The experiments were executed on a PC with a 2.50GHz CPU, and 512GB memory.

Table 2: Examined datasets.

Dataset	#Tuples	#Atts	Max vals per att
ACS	1,420,652	288	723
Stack Overflow	73,268	78	180
Flights	5,819,079	43	6952

Datasets, queries & claims. Similarly to previous work [33, 49, 66], we examined multiple common datasets and devised queries and claims inspired by real-world resources [6, 55].

<u>ACS</u>. We accessed the American Census Survey (ACS) data through the Folktables library [11]. We use the 2018 data that includes the seven largest US states: CA, TX, FL, NY, PA, IL, OH. This resulted in a dataset of 1,420,652 rows and 288 columns. Attribute names are encoded strings and attribute values are encoded numbers or strings. We mapped them to human readable strings using the ACS PUMS data dictionary.⁷ We also added two attributes based on groupings of the OCCP and NAICSP fields—the readable strings for these fields begin with a 3-letter encoding of the occupation field, which we extracted to a new attribute. The query is the average total income grouped by gender, and we search for refinements where women's income is higher than men's.

<u>Stack Overflow.</u> We used the Stack Overflow developer survey from 2022.⁸ The dataset contains 73K responses and 78 attributes, covering demographic, professional information, and the yearly compensation of participants. In some attributes, some rows contain multiple values, separated by a delimiter; in these, we kept only the first value. We focused on the median yearly compensation, and compared bachelor's degree graduates to master's degree graduates. Overall, the median salary of master's graduates was slightly higher, so we searched for refinements where the median salary of bachelor's graduates is higher.

Flights. The flight delays dataset⁹ contains 5.8M flight statistics (e.g., scheduled and actual departure and arrival times, carrier, airplane number) from 2015, over various carriers and airports. We joined the airports and flights relations to create a 43-attribute relation. We compared two weekdays, Monday and Saturday, and counted the flights with departure delays exceeding 10 minutes. On Mondays, there were more delays than on Saturdays (192,219 versus 134,681). We aim to endorse the claim of the reverse direction.

Data preprocessing. For numeric attributes other than the target attribute, we perform binning similarly to prior work [10, 44] The interval size is chosen according to the order of magnitude of the value range in the attribute. The start and end points of the bins are rounded to create natural-sounding ranges.

Algorithm variants. We examined the following prioritization methods for attribute combinations (Line 2 in Algorithm 1):

<u>MERGED TOP-k:</u> Interleaving merge combination of naturalness measure heuristics and pre-computations.

<u>SERIAL TOP-k:</u> Serial combination of naturalness measure heuristics and pre-computations. We iterate over the metrics from the

⁵https://github.com/shunita/claimendorse (accessed Oct 2024)

⁶https://huggingface.co/sentence-transformers/all-MiniLM-L6-v2 (accessed Oct 2024)

⁷https://www.census.gov/programs-surveys/acs/microdata/documentation.html (accessed Oct 2024)

⁸https://survey.stackoverflow.co/2022 (accessed Oct 2024)

⁹https://www.kaggle.com/datasets/usdot/flight-delays (accessed Oct 2024)

most accurately precomputed to the least accurate heuristic: *ANOVA*, *MI*, *EmbSim*, *StatSig*, and *Coverage*.

<u>*x*% SAMPLE:</u> The search is guided by a sample size of x% of the number of tuples in the database. We experimented with sample sizes of 1%, 5% and 10%. We first perform a search on the sample, and use the results to prioritize the attribute combinations in the search over the full database. The presented results for x% SAMPLE methods represent an average of 3 runs.

Baselines. We examine literature-based and naive baselines. Since the claim-endorsing problem is new, we compare our methods to similar problems through case studies (Section 5.2).

<u>Random order</u>: This baseline iterates over the attribute combinations in a random order. In the experiments below, the presented results for this baseline represent an average of 3 runs.

Single naturalness measure: This baseline prioritizes the attribute combinations according to a single naturalness measure. Each measure has a different prioritization method (detailed in Section 4.2.1).

<u>OREO [33, 34]</u>: OREO is designed to identify cherry-picked generalization statements by using a scoring function that considers all supporting subpopulations, weighted by size, where a higher score indicates stronger support. Additionally, OREO enables to discover counterexamples to the examined statement in the form of query refinements. As opposed to our approach, OREO provides *all* most general query refinements (as described in Section 3). This is compounded by the fact that OREO uses a predicate-level search instead of an attribute-level search (see Section 4.1), limiting scalability to high-dimensional datasets (up to 12 attributes in the examined datasets). Also, the refinements are *not ranked*.

For our case studies and user study, we use this method to find all maximal refinements. In our quantitative evaluation, we run an enhanced version of OREO with some differences: (1) Iterating over combinations of up to 2 attributes instead of all possible combinations (to handle large datasets, since as mentioned, original OREO does not scale to a large number of attributes); (2) Disabling generality filtering (Section 3) to retrieve a larger set of predicates; and (3) Anytime-style output - output the predicates as they are found instead of at the end of the run. We further give an unfair advantage to OREO by pre-selecting only the attributes responsible for top k predicates (found using other baselines). To avoid confusion, we name this version of OREO with the advantage as OREO^{*}.

<u>HypDB [48, 49]</u>: This system detects bias in average-group-by SQL queries. Given a query, HypDB finds a set of *covariates* (attributes with uneven distributions among the groups) that serve as an explanation for the query results, based on causal analysis. Each selected attribute is associated with a responsibility score. Although this approach does not directly offer query refinements supporting the user claim, it highlights attributes with uneven distributions among the groups defined by the grouping attribute. However, we demonstrate that attributes relevant to defining natural refinements that support the user's claim are not necessarily covariates. In our effectiveness evaluation, we use the responsibility score to prioritize the attribute combinations, where the score of an attribute combination is the sum of responsibility scores of each attribute.

Table 3: Example refinements found through MERGED TOP-k.

Dataset	Predicate	Avg. Nat	Rank	Time (s)	
0. 1	OpSysProfUse=Linux-based	0.44	32	2	
Stack	\land YearsCode=0-10				
Overflow	Employment = Full-time∧	0.10	59 (Cov.)	13	
	OrgSize = "1K-5K"	0.10	37 (007.)		
ACS	GradeLevelAttending= $12 \land$	0.40	73	197	
	Occupation=Customer Service	0.40	73		
	Occupation=Cooks ∧	0.41	20	70	
	HoursPerWeek=10-20	0.41	20	70	
	MaritalStatus=NeverMarried	0.39	56	475	
	∧ WhenLastWorked=>5y ago	0.39	20	4/5	
Flights	Airline=Hawaiian Airlines Inc.	0.36	1	36	
riignts	ScheduledDeparture=3:00-4:00	0.30	2	40	

Evaluation metric. We report the *score recall* of each examined baseline. For a generated set r of refinements, and naturalness measure v, let v_i^r denote the top i'th v score found in r. Define $S_r := \sum_{i=1}^k v_i^r$. The *score recall* of r is given by: S_r/S_{full} where S_{full} is sum of top-k scores of an exhaustive search. While there are other metrics to evaluate retrieval results, they are not suitable for our task. Classic recall (based on exact matches), Normalized Discounted Cumulative Gain (NDCG) [21], and Kendall's- τ [29], are all based on the existence of an underlying ground truth ranking. However, this is not the case here, as there can be many results of similar quality.

Default configuration. We use a default of m=2 maximal number of atoms in each combination. Even with this limitation, predicate search is computationally intensive. For example, in ACS, there are 120 split attributes, and 7140 2-attribute combinations, resulting in 7260 pairwise attribute combinations. Processing all 7260 takes over 1.5 hours, thus creating an experimental setup where the differences between methods can be evaluated. As default we use k=100.

5.2 Case Studies

We use MERGED TOP-*k* prioritization, which achieved the best results among our methods in terms (see Sections 5.4 and 5.5 in the sequel). We note that the choice of prioritization method affects the time it takes to find the best refinements, but if given unlimited time, all prioritization methods retrieve the same set of refinements; the prioritization only affects the order in which the attribute combinations are searched, but eventually, we iterate over all of them. In this section, we allow the search to finish and compare the final set of retrieved refinements to those achieved by existing solutions, since existing solutions have no time limit. Nevertheless, our approach retrieves convincing refinements that endorse the claim in reasonable time (Table 3). We compare MERGED TOP-*k* with existing solutions on three example scenarios, to showcase the distinctions in our results. Specifically, we consider OREO and HypDB.

Stack Overflow. We consider the claim "the average salary of people with M.Sc. is higher than that of people with B.Sc."¹⁰ MERGED TOP-*k* discovered meaningful and comprehensible refinements that endorse the claim, as shown in Table 3. One of the top scoring refinements is (OpSysProfessional use = Linux-based) \land (YearsCode = 0 - 10) (Avg. nat.=0.44, ranked 32). The rationale is that Hi-Tech workers who use Linux are typically those in technical positions, and for those with little experience, a master's degree may well contribute to a higher salary. Ranking by different measures, we get additional results. With *Coverage*, a top refinement is (Employment =

 $^{^{10}\}mathrm{We}$ used Average since HypDB is suitable only for average aggregation.

Full-time) \land (Org. size = "1K-5K") that covers over 3000 people. Both predicates were found in under 13 seconds.

Due to the small size of the dataset, and the limitation of two atoms, OREO returned a manageable set of 282 maximal predicates. Some are quite convincing, like DevType = "data scientist or ML specialist". In this case, it may be beneficial to combine the two approaches, by incorporating OREO's output in the refinements we rank. However, the results show that being a maximal predicate may not fully capture our objective. Maximal refinements may reflect a side effect of the survey design, and not necessarily a characteristic of the group, like Ethnicity="I don't know" (Avg. nat.=0.23, ranked 2785). Further, the set of maximal predicates may omit convincing refinements, e.g., the above-mentioned (OpSysProfessional use =Linux-based) \land (YearsCode=0 - 10), which is statistically significant and shows a considerable difference in salaries (\$120.1K vs. \$133.6K), because (OpSysProfessional use = Linux-based) satisfies the claim with a tiny difference (\$165.7K vs. \$167.2K).

For explaining the impact of M.Sc. versus B.Sc. on the average salary, the covariates (and responsibility scores) found by HypDB are BuyNewTool (0.34), LanguageWantToWorkWith (0.33), and RemoteWork (0.33). These attributes accounted for 8 out of the 224 single-atom refinements that satisfy the claim, while RemoteWork did not yield any predicates. The average naturalness score of these attributes (i.e., their corresponding predicates) ranges between 0.04 - 0.19. Our viewpoint concurs as they are not very intuitive for supporting claims. Other predicates that satisfy the claim had a much higher average naturalness score.

ACS. We consider the claim "The average salary for women is higher than the the average salary for men". MERGED TOP-k found meaningful and understandable refinements that endorse the claim, as shown in Table 3. For example, there are 27 occupations where women's average income is higher, e.g., medical transcriptionists and tutors. Other examples are high school students attending the 12th grade, and people living with a single working parent.

OREO identified 2,646 maximal predicates supporting this claim. This is a larger number of predicates than the user can usually consider. Among these, OREO prioritizes the 97 single-atom refinements but lacks a method to distinguish between predicates or determine their naturalness. For instance, OREO does not differentiate between the predicates: "People who have Indian Health Service" and "People who have a doctorate degree".

The top-3 covariates (and scores) by HypDB were "Gave birth within past year" (0.27), "Veteran Service Disability Rating" (0.27), and "Year of naturalization" (0.25). Out of the 6 covariants, only one ("Veteran Service Disability Rating") yielded any legal refinements. The refinement is defined by "Veteran Service Disability Rating"="Not reported," which contains a small group of women (58) and a small difference between the average incomes of men and women. The found covariates are attributes that can explain income gaps between the genders. For example, giving birth in the past year may harm the yearly income, due to unpaid leave from work. However, as demonstrated, a covariate attribute does not necessarily define a subgroup where the opposite direction holds.

Flights. We consider the claim "Departure delays of more than 10 minutes are more common on Saturdays than Mondays." Here we use m = 1. The difference between the approaches is similar to the

previous cases. Some refinements obtained from MERGED TOP-k are in Table 3. MERGED TOP-k returned 2,300 refinements that endorse the claim. The refinement with the highest average naturalness score (0.36) was Hawaiian Airlines. Another refinement is flights scheduled to depart between 3AM and 4AM. However, for the opposite claim (departure delays are more common on Mondays), the algorithm returned many more refinements (6,488) and the highest average naturalness was 0.57, distinctively higher than the highest naturalness for the opposite claim (0.36). This case shows the merit of using our framework as a touchstone for verifying claims, in addition to endorsing a given claim through refinements.

The use cases demonstrate that claim endorsement cannot be directly solved by finding covariates of an intervention on an outcome, as covariate attributes do not necessarily yield successful refinements that endorse the claim. Furthermore, sorting the naturalness of refinements is one of the pillars of this work, but it is not discussed in HypDB and OREO. Finally, OREO employs a predicatelevel search, and as we show in the full version [2], our attribute level approach is faster by an order of magnitude. Nevertheless, we compare our approach to these baselines in the next experiments.

5.3 User Study

We conducted a user study (1) to evaluate the extent to which the naturalness measures correspond to intuitive naturalness, (2) to find out which naturalness measures are preferred, (3) to inspect the effect of generality filtering adapted from OREO, and (4) to compare the measures to the HypDB responsibility score.

For each dataset (and corresponding claim), we generated several statements supporting the claim¹¹ We presented the statements to 50 users through the Prolific Academic platform¹² and asked them to rate, on a scale of 1 to 5, how much they would recommend using each to an author of an article about the claim. We included the statements with the maximal score in each measure, along with the statement with the maximal HypDB responsibility. For each of these, we also applied a generality filter (Section 3) and chose the statement with the maximal score out of the most general statements, thus combining our measures with the generality property of counter arguments used in OREO [34]. We also retrieved the statements with the median score in each measure. For each choice method we computed the average rating, and the number of times its statements were ranked at the top of their group (Figure 2).

We conclude the following. (1) The refinements in the medians of the measures had significantly lower ratings than their maximum counterparts, hence the naturalness measures coincide with naturalness as perceived by the participants. (2) Maximal *Coverage* was marked best more times than any other method, followed closely by maximal *StatSig*. The highest average rating belongs to maximal average naturalness with generality filtering. (3) Generality filtering, adapted from OREO [34] and applied on each selection method, increased the rating of each method, and statistically significantly for *StatSig*, *EmbSim* and average naturalness. (4) HypDB responsibility score was significantly lower than max *Coverage*, max *StatSig* and max average naturalness, which coincides with our observations in our case studies, that HypDB solves a different problem.

¹¹The full set of survey claims and statements is included in the full version [2].
¹²https://www.prolific.com (accessed Oct 2024)

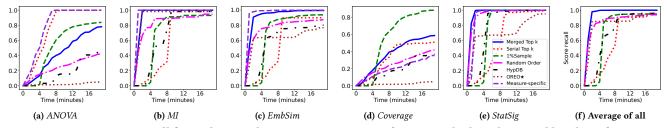


Figure 1: Top 100 score recall for each naturalness measure over time for our methods and external baselines for ACS.

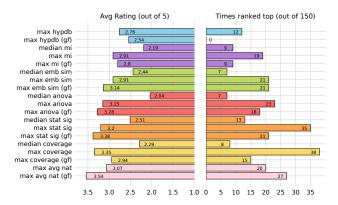


Figure 2: User study results: average rating and times ranked top for each method. (gf) stands for generality filter.

5.4 Effectiveness Evaluation

We compare our methods and external baselines based on score recall over time. Table 4 shows the time till score recall 95% for each prioritization method and naturalness measure. The score recall over time for each naturalness measure for ACS is shown in Figure 1. We focus here on the ACS dataset since it has the largest number of attributes and over 1M rows, and therefore requires the longest run times. Similar trends were seen for the other datasets.

For most of the naturalness measures and datasets, the best method was MERGED TOP-k. In ACS, this method was the first to reach 0.95 recall for MI, EmbSim, StatSig, and the average of all naturalness measures. Over all naturalness measures and all datasets, MERGED TOP-k was on average 13.4 (and up to 78.3) times faster to reach 0.95 recall than the random-order baseline, and on average 16.2 (up to 82.5) times faster than HypDB. For the Flights dataset, SERIAL TOP-k was the fastest to reach 0.95 recall for all measures except Coverage, with MERGED TOP-k as a close second. The best prioritization for ANOVA was SERIAL TOP-k, simply because it is the first measure in the serial order. The external baselines yielded inferior results for all datasets and measures. Prioritizing by HypDB scores performed similarly to random. OREO* was not applicable to the median query used in stack overflow (since median is not supported in OREO), but on ACS and Flights it was slower to reach 95% recall, if it reached it at all. This is likely because even the enhanced OREO* version we used operates on a predicate level, which is much more time consuming than our attribute level solution.

The *Coverage* score recall was the slowest to reach 95% prioritization methods (e.g., over 48 minutes for ACS). This is because *Coverage* is a predicate-level measure, and the hardest to provide an accurate heuristic for. Our heuristic of prioritizing attribute combinations by their number of large groups hardly predicted large groups that satisfy the user. Still, in Stack Overflow, MERGED TOP-*k* was the fastest to reach 95% score recall for *Coverage*. For the ACS dataset, the sampling method with 1% was the fastest, despite the long preprocessing time. We conclude that in the absence of an accurate heuristic for a given measure, sampling can yield fast results, but for naturalness measures with a good predictor (heuristic or pre-computation), SERIAL TOP-*k* and MERGED TOP-*k* prevail.

For the *x*% SAMPLE method we experimented with sample sizes of 1%, 5%, and 10%. The complete set of results is in the full version [2]. We observed a trade-off between the time until the first result (which grows with the sample size) and the time until a high score recall (which is shorter for larger samples, as the preliminary search yields more accurate results). However, for most naturalness measures, 1%, so we consider it a recommended sample size. Similar trends were seen for the other datasets, as shown in Table 4.

Due to the results of x% SAMPLE on the *Coverage* measure, one may consider to integrate the sampling method into the MERGED TOP-*k* or SERIAL TOP-*k* methods, as another ranking of the attribute combinations. However, the pre-processing of x% SAMPLE will increase the time until high recall is reached in all other measures.

The anytime nature of the algorithm is reflected in that the set of retrieved refinements grows, thus the sum of top-k scores monotonically increases, as seen in Figure 1. In the following experiments, we used this observation to optimize the experimentation time by stopping when the score recall reaches a critical threshold, computed with reference to a full run of the algorithm. When the score recall is not known, one could design an early stopping strategy, based on the size of the change in the top k sum of naturalness scores. For example, "If the c last attribute combinations did not improve the sum of top-k v scores by more than ϵ , stop the search".

5.5 Parameter Sensitivity

Sensitivity to number of tuples and columns. For the number of tuples (Figure 3a), we sample varying amounts of tuples out of the full ACS dataset. For the number of columns (Figure 3a), we sample increasing sizes of subsets of SAtt(D) in the ACS dataset. For each number of tuples or columns, we show the average of three runs. We measure the time until a 95% score recall was achieved, for the naturalness measures average score, in each of our five main prioritization methods and in the external baselines - HypDB and OREO^{*}. Similar trends were observed for the other two datasets.

For the naive prioritization of random order and for HypDB, the time until 95% recall steadily rises with the number of tuples or columns. This is also the case for 1% SAMPLE, since the time to run

Table 4: Time (seconds) to 0.95 score recall and first result time (FRT) for different prioritizations and baselines (OREO* and HypDB). OREO* is inapplicable to Stack Overflow as it does not support median aggregation. Entries "t/o" mean that timeout is reached (2 hours for ACS7 and 5 hours for Flights).

Dataset	Prioritization	EmbSim	StatSig	ANOVA	MI	Coverage	Avg.	FRT
	1% Sample (3)	1303	300.7	3881	833.3	2886	401	191.0
$\begin{array}{l} ACS\\ \alpha = Avg \end{array}$	5% SAMPLE (3)	734	598.7	5733.3	584.33	3720.7	588.3	582.4
	10% SAMPLE (3)	1006.3	839.3	3812	830.3	3895.7	834	828.1
	Rand. Order (3)	2914.3	177	5129	909.7	5518.7	1479	10.2
	Serial Top- k	3415	554	469	503	4566	1218	84.3
	Merged Top- k	231	71	1557	66	4886	76	49.9
	HypDB	1992	290	6079	502	4272	722	139.9
	OREO*	5913	1051	t/o	98	t/o	982	38.5
	1% Sample (3)	63.3	41.7	298	43.3	332.3	40.7	38.1
	5% SAMPLE (3)	77.3	57.7	100.3	54.7	358	55	53.5
Stack O.	10% Sample (3)	94.3	71.7	99.3	70	362.7	70	68.7
$\alpha = Med$	Rand. Order (3)	206.3	9	207.3	313.3	290.7	200	1.1
u = meu	Serial Top- k	115	9	4	5	246	7	1.8
	Merged Top- k	25	7	11	4	206	7	1.9
	HypDB	280	28	313	330	261	203	9.7
	1% Sample (3)	133.7	-	131.3	131.3	1022.3	131.3	129.8
	5% SAMPLE (3)	261	-	259	259	1148.7	259	257.4
El: dete	10% SAMPLE (3)	344	-	342	342	1232.7	342	340.4
Flights $\alpha = Cnt$	Rand. Order (3)	111.7	-	512.7	381.3	793	435.3	20.9
$\alpha = Cnt$	Serial Top- k	37	-	35	35	895	35	34.5
	Merged Top- k	40	-	38	38	932	38	34.6
	HypDB	239	-	979	979	928	979	89.9
	OREO*	13050	-	t/o	t/o	12298	t/o	12.7

the pre-processing step of the search on the 1% sample naturally depends on the number of tuples in the database. For OREO \star , the time to reach 95% recall was on average 7.8X longer than random order for the number of tuples experiment, and 84X longer for the number of columns experiment. This is due to the predicate level approach taken in OREO*, which does not scale to large datasets. However, for MERGED TOP-k, the time until 95% recall is almost constant. This is due to the prioritization of the search, and covering the most promising attribute combinations at the beginning. For SERIAL TOP-k, there is a rise in the time until 95% score recall with the number of tuples or columns, although not as steep as the random order baseline. A possible explanation is that in SERIAL TOP-k, only after k refinements for a specific naturalness measure are found, do we continue to the next measure. If there is only a small number of refinements for a specific measure, this may delay the recall for the other measures (and for their average).

Sensitivity to k (number of top refinements). We measured the time until 95% top-k score recall of the average naturalness measures in the five prioritizations with k values from 100 to 1000 on ACS (Figure 3c). Similar trends were observed on the other datasets. For all methods except SERIAL TOP-k, the choice of k does not affect the search algorithm, only the recall computation. For SERIAL TOP-k, k refinements are fetched from each naturalness ranking at the beginning of the search. Therefore, for SERIAL TOP-k each point in Figure 3c represents a different algorithm run.

As expected, for all methods the time to 95% recall grows with k. However, MERGED TOP-k exhibited the most subtle rise in time to 95% recall out of all methods (ranging from 1.3 to 3.6 minutes), showing the merit of this combined prioritization method.

Sensitivity to *m* (maximal number of attributes per combination). Increasing *m* may lead to complex statements, which are less natural. For example, (OpSysProfUse=Windows) \land (YearsCode = 0-10) \land (OrgSize = 20-99) \land (DatabaseHaveWorkedWith = MariaDB) \land (AccessibilityDifficulties = None). Most refinements with a large *m* are specializations of a more general refinement (e.g., for stack overflow with 10 attributes, 99.2% of the 44,890 refinements with 5 atoms specialize 3-atom refinements). While the run times of our algorithms are relatively short, the precomputation times for computing *ANOVA* and *MI* were very long (about 24 hours for *m*=3). Therefore, we limited this experiment to the 10 attributes responsible for the largest number of refinements retrieved in the 2-atom search. Due to OREO*'s long run time, it was not included in this experiment. The results are shown in Figure 3d.

As previously, MERGED TOP-k and SERIAL TOP-k were the fastest to reach 95% recall. For comparison, we added the time to conduct a full claim endorsement search (regardless of the recall level), and as expected, it grows exponentially with m. It can be seen that MERGED TOP-k and SERIAL TOP-k were much faster than that.

6 RELATED WORK

Fact-checking and cherry-picking. Automatic methods to identify cherry-picked results and fake news have been widely studied [17, 20, 68]. Traditional fact-checking methods rely on domain knowledge [18, 19] and lack scalability and persuasiveness without supporting datasets. Other methods employ machine learning and NLP techniques for efficient computational fact-checking [15, 17]. A considerable body of research focuses on automating fact-checking using structured data [3, 22, 33, 62, 63]. In one line of work [62, 63], parameterized queries are defined, and the robustness of a fact to parameter perturbations is analyzed. Perturbations are assessed for their relevance and naturalness, depending on the attributes and domain knowledge. In another line of work [3, 4], trendlines are examined with respect to their robustness to changes in their start and end points. Cherry-picking has also been examined in rankings [4], where items are ranked according to a linear combination of numeric attributes, and the weights can be cherry-picked.

Explanations of query answers. Claim endorsement can be viewed as the opposite of the intervention or responsibility approaches in query result explanations [37-39, 46, 47, 60]. In the latter, tuples conforming to a predicate are removed from the database so that the result on the smaller database satisfies an assertion. Conversely, a predicate in our setting qualifies the tuples that remain in the database to satisfy a claim. A commonly used explanation type for query results is a set of predicates that differentiate between tuples in the answer of aggregate queries [14, 31, 46, 47, 53, 54]. Our framework similarly provides a ranked list of predicates that define the refinements where the claim holds. However, our primary goal differs in that we seek the most natural refinements to endorse a claim, rather than predicates explaining unexpected answers. Query refinements have also been used for improving query results according to desired properties [8], including the number of results [9, 30, 41-43, 57], diversity [32], or bias removal [49].

Multidimensional data aggregation. Previous work on multidimensional data aggregation has extended the traditional drill-down and roll-up operators to find interesting data parts for exploration [1, 23, 50, 65]. Others works have focused on assessing the similarity between data cubes [5]). None of these methods corroborate a

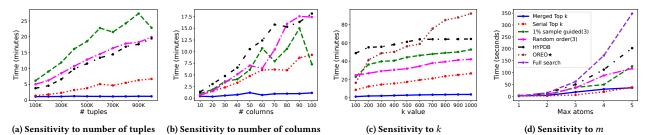


Figure 3: Time for 95% score recall of avg. naturalness over ACS with at most m = 2 atoms, with varying number of (a) tuples, (b) columns, and (c) varying values of k. (d) Sensitivity to maximal number of atoms (m): time for 95% recall of average naturalness with increasing number of atoms over Stack Overflow with 10 attributes.

user's claim in the aggregate view, or check if it represents a meaningful definition of a subpopulation for the claim. Some employ the CUBE operator [1], which may seem relevant here to materialize all combinations of attributes for refinements. Yet, this approach is not scalable; database systems usually limit the number of attributes in the cube operator to 12 [49], due to its size being exponential in the number of attributes. It has been shown [33] that over 6 attributes, CUBE is impractical for its time and memory consumption.

View recommendation. Another relevant area of research has used data aggregations to discover intriguing data visualizations [10, 56], with various techniques proposed to recommend the top-k most interesting views [12, 13, 36]. Some studies recommend visualizations based on visual quality functions, such as visual expressiveness [59]. Others use deviation-based functions, where the interestingness of a view is measured by its distance from a reference view [10, 12, 56]. Additional studies have explored multi-objective utility functions that capture different aspects of interestingness (e.g., visual quality, deviation) or aimed to discover the most appropriate utility function for the current analysis context [13, 36, 67]. The key distinction from the current work is that the interestingness differs significantly from the naturalness; in fact, they can be seen as almost opposite concepts. While they seek to identify interesting views that reveal anomalies in the data, our objective is to find the most natural views-those with the fewest special characteristics that do not attract particular attention. The goal is to support claims that hold true across natural subpopulations.

Nonetheless, we share common techniques with this line of work. We employ sharing-based optimizations to minimize the number of query executions, similarly to [56], and pruning-based optimizations (like filtering attribute combinations when their maximum group size is too small) to improve running times, as done in [56, 59].

7 CONCLUSIONS AND FUTURE DIRECTIONS

We presented a framework for endorsing user claims through query refinements. Our framework quantifies the refinement quality using naturalness measures adapted from the literature but can support any naturalness measure. We proposed an efficient approach for computing refinements using an anytime algorithm.

There are multiple avenues of future work. First, it is possible to extend our work to a wider problem definition. Our work currently considers a single relation and can be extended to multi-relation databases, by creating a single view in advance, or by modifying the query for each attribute combination to enable multiple relations. While we focused on refinements (assuming that the claim does not hold in general), ideas in this paper could also be used for relaxations. Attribute combinations can be created from attributes appearing in the where clause of the query. These combinations can be prioritized as in Section 4.2 and searched as in Section 4.1. Relaxation for conjunctions appears to be a simpler problem, with the main challenge being prioritization, where our methods also apply. An important direction is to enrich the predicate language to inequalities and disjunctions. Inequality predicates can easily be supported in the algorithmic framework, while disjunctions would require changes to the algorithm and different optimizations.

Second, one may consider methods to make the results more accessible to users. For example, one can incorporate desiderata for the ranking of refinements that considers the entire list, such as diversification which would introduce variation in the refinements. This would require designing metrics that combine both naturalness and diversity, and re-ranking the returned refinements while considering the new metric. Alternatively, one can consider a summarization strategy for results, grouping similar refinements and thus facilitating the analysis process.

Third, to make the search process interactive, decision rules could guide users on when to stop the search, e.g., based on result quality bounds. Another intriguing future work is to discover the most suitable naturalness measure in the current context based on user interaction (inspired by [67]).

Finally, our work considers user claims that compare two groups. We intend to consider more elaborate claims such as trends in the query results. This may require considerable adjustment of the problem definition, the predicate space and the algorithm.

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