

General Database Statistics Using Entropy Maximization

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Abstract. We propose a framework in which query sizes can be estimated from arbitrary statistical assertions on the data. In its most general form, a statistical assertion states that the size of the output of a conjunctive query over the data is a given number. A very simple example is a histogram, which makes assertions about the sizes of the output of several range queries. Our model also allows much more complex assertions that include joins and projections. To model such complex statistical assertions we propose to use the Entropy-Maximization (EM) probability distribution. In this model any set of statistics that is consistent has a precise semantics, and every query has an precise size estimate. We show that several classes of statistics can be solved in closed form.

1 Introduction

Modern database query optimizers are the result of thousands of man years worth of work from very talented individuals, and so are extremely sophisticated. Although the optimizers themselves are sophisticated, the heuristics they employ are often not: When estimating the selectivity of two predicates, the optimizer might make an *independence assumption* and so, assume that the selectivities of two predicates are independent. When estimating the size of the intersection of two columns, the optimizer might make a *containment assumption* and assume that the values in one column are contained in the other. The cleverness of the optimizer is in *how* and *when* to apply these rules. In spite of this intense effort and the fundamental importance of the problem, there is no general theory that explains how the optimizer should make these choices or when such choices are consistent. In this paper, we take a first step towards such a general, principled theory of how optimizers should make use of statistics.

The lack of a principled framework is likely to become an even more critical problem as new sources of statistics become available to the query optimizer. For example, several proposals [3, 17] advocate acquiring *query feedback* and incorporating this statistical feedback into the optimizer. It is easy to collect cardinality statistics from each query as it is executed by the database engine. The difficulty lies in combining this newfound plethora of statistics to produce a single, principled estimate. The lack of such a principled framework is a major reason that execution feedback has not been widely adopted in commercial database engines.

The key object of our study is a *statistical program*, which is a set of pairs (v, d) , where v is a query (also called a view) and $d > 0$ is a number. Each pair

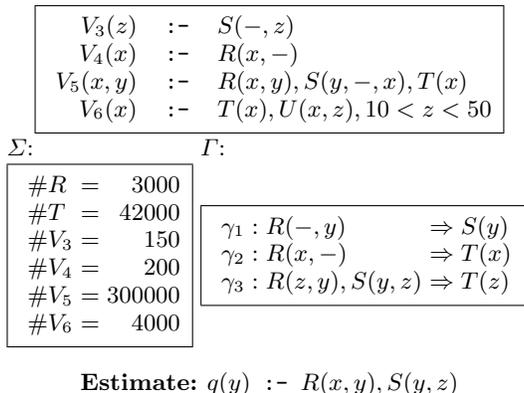


Fig. 1. An example of a Statistical Program and a query, q whose cardinality we would like to estimate.

(v, d) is called a *statistical assertion*, and means intuitively that the answer to v has expected size d ; we write it as $\#v = d$ and say simply that “the size of v is d ”. A statistical program encodes the information that is available to an optimizer. The primary use of this information is to estimate the expected result size of other queries during query optimization.

Example 1. Figure 1 illustrates a statistical program that asserts the sizes of 4 views and 2 base relations. The program also asks for the estimated size of another query. This program asserts that the expected number of distinct tuples in the relation R is 3000. The program also asserts (via V_3) that the second attribute of S contains 150 values. Our proposal also allows complex assertions that involve joins and arithmetic predicates, such as V_6 . In addition to statistical assertions, our model also allows specifies *inclusion constraints*. These constraints are *hard constraints* and must hold in *every instance* I the distribution considers possible, i.e., for which $\mathbf{P}[I] > 0$. For example, γ_1 says that each value in the second column of R is also in the S relation.

In this paper, we study the probability distribution over database instances that satisfy a given statistical program Σ .

1.1 Our Approach: Entropy Maximization

In this paper we define a model for a statistical program using the Entropy Maximization principle. We assume that the relations in the database are drawn from a finite domain D , of size N , and that each database instance I has some probability $\mathbf{P}(I)$. \mathbf{P} is chosen to fit the statistics Σ and without making any other assumptions on the data. More precisely (1) for each assertion $\#v = d$ in Σ , the expected size of v under \mathbf{P} is d , and (2) the probability distribution \mathbf{P} has the maximum entropy among those that satisfy (1) (formal definition given in Sec. 2). The EM model for a statistical program Σ is an instance of the

general EM principle in probability theory, discussed for example by Jaynes [12, Ch.9,11], and which has also been applied to consistent use and construction of histograms [14, 16].

The EM framework has many attractive features. First, any combination of statistical assertions has a well-defined semantics (except, of course, when it is inconsistent); thus, a statistical program is treated as a whole, as opposed to a set of separate synopses. Second, every query has a well defined cardinality estimate; there is no restriction on the query, and the query estimate no longer depends on which heuristics are used to do the estimation. A third reason is that the EM framework has an interesting property that allows us to add a new statistical assertion smoothly: if the estimate of a query q under a statistical program Σ is d , then after adding the assertion $\#q = d$ the new EM probability distribution is identical to the previous one. In practice this means that if we add a small correction to the model, $\#q = d'$ where $d' \approx d$, then the model will change smoothly. The final, and most important conceptual reason is that in a precise sense the probability distribution given by entropy maximization depends *only on the provided statistics* and makes no additional assumptions beyond this. We return to this point when we formally define our model and state its properties in Sec. 2.1.

In this paper, we study the following *model computation problem*: given a statistical program Σ and a set of full inclusion constraints Γ , find a solutions to the EM model. (We explain below the reason for introducing constraints.) Since an EM solution is tied to the particular domain D , we seek to remove the dependency by letting the domain size N grow to infinity as is done, for example in random graphs [9], knowledge representation [2], or asymptotic query probability [5]. Since we seek an analytic understanding of the model, our goal is to find analytic (asymptotic) solutions to the EM model. Solving the EM model in general is, however, a very hard problem. We report in this paper several partial results on the asymptotic solutions for statistical programs. While our results do not add up to a comprehensive solution to statistical programs they do offer explicit solutions in several cases, and shed light on the nature of the EM model for database statistics.

1.2 Main Technical Results

In this paper, we introduce classify programs according to two axes. First, whether the statistical assertions are on base tables only, or on both base tables and views: we say that Σ is in *normal form* (NF) if all statistical assertions are on base tables; otherwise it is in non-NF. The program in Fig 1 is in non-NF. Second, whether the views in Σ, Γ have joins or are join-free: we call the program *composite* if all views have joins, and *atomic* if all are join-free (we do not consider mixed atomic/composite programs). All four combinations are possible, e.g., an NF, composite program means that all statistical assertions are on base tables and the inclusion constraints are composite views. In this paper, we consider only *composite programs*³.

³ It turns out that atomic and mixed programs require an entirely different set of more complex, analytic techniques. In the full version of this paper [13], we discuss our

We give a complete solution for composite, NF programs: given statistics on the base tables and a set of full inclusion constraints, the EM model is described by an explicit formula. We prove this by using techniques from [6]. Second, we prove a more limited result for non-NF programs, by giving an explicit formula when the views are restricted to *project-semi-joins*. This explicit formula gives an important insight into the nature of difficulty of the non-NF programs, as we explain below.

In addition to these explicit solutions, we discuss a generic technique that we encapsulate as the *conditioning theorem* (Sec. 3.1); this reduces a more complex program to a simpler program plus additional inclusion constraints; this is what has motivated us to study statistical programs *together* with constraints.

The conditioning theorem states that every solution \mathbf{P} to the EM model can be expressed as a conditional probability $\mathbf{P}(-) = \mathbf{P}_0(- \mid \Gamma)$, where \mathbf{P}_0 is a tuple-independent distribution called the *prior*, and Γ is a set of inclusion and set-equality constraints. Since \mathbf{P}_0 is tuple-independent, it is specified by a number $p_i \in (0, 1)$, one for each relation R_i , representing the probability that a generic tuple in the domain belongs to R_i ; the corresponding *odds* is denoted $\alpha_i = p_i/(1 - p_i)$. Understanding the quantity $\mathbf{P}_0(\Gamma)$ is a key component to understanding the EM model. This quantity converges to 1 for composite NF programs, and to 0 for composite NNF programs, suggesting that the techniques used to solve composite, NF programs (which turn out to be simpler) do not extend to the other cases.

The EM model computation that we study in this paper is the first step in our program of using the EM framework for query size estimation. The second step is to use the model in order to do query size estimation, which we leave for future work, noting that it has been solved for the special case of independent distributions [6].

The rest of the paper is organized as follows. We describe the EM model in Sec. 2, discuss composite programs in Sec. 3, and discuss how to handle range predicates in Sec. 4. We conclude in Sec. 6.

2 The EM Model

We introduce basic notations then review the EM model. \mathbf{CQ} denotes the class of conjunctive queries over a relational schema R_1, \dots, R_m . We define a *project-join* query as a conjunctive query without constants and where no subgoal has repeated variables, and write \mathbf{PJ} for the class of project-join queries. For example $R(x, y), S(y, z, u)$ is a project-join query, but neither $R(a, x)$ nor $S(x, x, y), T(y, z)$ are. An arithmetic predicate, or range predicate, has the form $x \text{ op } c$, where $\text{op} \in \{<, \leq, >, \geq\}$ and c is a constant; we denote by \mathbf{PJ}^{\leq} the set of project-join queries with range predicates.

Let Γ be a set of *full inclusion constraints*, i.e., statements of the form $\forall \bar{x}. w(\bar{x}) \Rightarrow R_i(\bar{x})$, where $w \in \mathbf{PJ}^{\leq}$ and R_i is a relation name.

preliminary results for atomic programs. We show for example that *any* program can be transformed to a normal form program. Interestingly, this normalization process also introduces additional inclusion constraints.

2.1 Background: The EM Model

For a fixed domain D and constraints Γ we denote $\mathcal{I}(\Gamma)$ the set of all instances over D that satisfy Γ ; the set of all instances over D is $\mathcal{I}(\emptyset)$, which we abbreviate \mathcal{I} . A probability distribution on $\mathcal{I}(\Gamma)$ is a set of numbers $\bar{p} = (p_I)_{I \in \mathcal{I}(\Gamma)}$ in $[0, 1]$ that sum up to 1. We use the notations p_I and $\mathbf{P}[I]$ interchangeably in this paper.

A *statistical program* is a pair $\Sigma = (\bar{v}, \bar{d})$, where $\bar{v} = (v_1, \dots, v_s)$ are project-join queries, $v_i \in \text{PJ}$, and (d_1, \dots, d_s) are positive real numbers. A pair (v_i, d_i) is a *statistical assertion* that we write informally as $\#v_i = d_i$; in the simplest case it can just assert the cardinality of a relation, $\#R_i = d_i$. A probability distribution $\mathcal{I}(\Gamma)$ *satisfies* a statistical program Σ if $E[|v_i|] = d_i$, for all $i = 1, m$. Here $E[|v_i|]$ denotes the expected value of the size of the view v_i , i.e., $\sum_{I \in \mathcal{I}} |v_i(I)| p_I$. We will also allow the domain size N to grow to infinity. For fixed values \bar{d} we say that a sequence of probability distributions $(\bar{p}_N)_{N > 0}$ *satisfies* $\Sigma = (\bar{v}, \bar{d})$ *asymptotically* if $\lim_{N \rightarrow \infty} E_N[|v_i|] = d_i$, for $i = 1, m$.

Given a program Σ , we want to determine the most “natural” probability distribution \bar{p} that satisfies Σ and use it to estimate query cardinalities. In general, there may not exist any probability distribution that satisfies Σ ; in this case, we say that Σ is unsatisfiable. On the other hand, there may exist many solutions. To choose a canonical one, we apply the Entropy Maximization (EM) principle.

Definition 1. A probability distribution $\bar{p} = (p_I)_{I \in \mathcal{I}(\Gamma)}$ is an EM distribution associated to Σ if the following two conditions hold: (1) \bar{p} satisfies Σ , and (2) it has the maximum entropy among all distributions that satisfy Σ , where the entropy is $H = -\sum_{I \in \mathcal{I}(\Gamma)} p_I \log p_I$.

With slight abuse, we refer to an EM distribution as *the* EM model, assuming it is unique. For a simple illustration, consider the following program on the relation $R(A, B, C)$: $\#R = 200$, $\#R.A = 40$, $\#R.B = 30$, $\#R.C = 20$. Thus, we know the cardinality of R , and the number of distinct values of each of the attributes A, B, C . We want to estimate $\#R.AB$, i.e., the number of distinct values of pairs AB . Clearly this number can be anywhere between 40 and 200, but currently there does not exist a principled approach for query optimizers to estimate the number of distinct pairs AB from the other four statistics. The EM model gives such a principled approach. According to this model, R is a random instance over a large domain D of size N , according to a probability distribution described by the probabilities p_I , for $I \subseteq D^3$. The distribution p_I is defined precisely: it satisfies the four statistical assertions above, and is such that the entropy is maximized. Therefore, the estimate we seek also has a well defined semantics, as $E[\#R.AB] = \sum_{I \subseteq D^3} p_I |I.AB|$. This estimate will certainly be between 40 and 200; it will depend on N , which is an undesirable property, but a sensible thing to do is to let N grow to infinity, and compute the limit of $E[\#R.AB]$. Thus, the EM model offers a principled and uniform approach to query size estimation. Of course, in order to compute any estimate we must first find the EM distribution p_I ; this is the goal in this paper.

To describe the general form of an EM distribution, we need some definitions. Fix the set of constraints Γ and the views $\bar{v} = (v_1, \dots, v_s)$.

Definition 2. The partition function for Γ and \bar{v} is the following polynomial T with s variables $\bar{x} = (x_1, \dots, x_s)$:

$$T^{\Gamma, \bar{v}}(\bar{x}) = \sum_{I \in \mathcal{I}(\Gamma)} x_1^{|v_1(I)|} \dots x_s^{|v_s(I)|}$$

Let $\bar{\alpha} = (\alpha_1, \dots, \alpha_s)$ be s positive real numbers. The probability distribution associated to $(\Gamma, \bar{v}, \bar{\alpha})$ is:

$$p_I = \omega \alpha_1^{|v_1(I)|} \dots \alpha_s^{|v_s(I)|} \quad (1)$$

where $\omega = 1/T^{\Gamma, \bar{v}}(\bar{\alpha})$.

We write T instead of $T^{\Gamma, \bar{v}}$ when Γ, \bar{v} are clear from the context. The partition function can be written more compactly as:

$$T(\bar{x}) = \sum_{k_1, \dots, k_s} C_{\Gamma}(N, k_1, \dots, k_s) x_1^{k_1} \dots x_s^{k_s}$$

where $C_{\Gamma}(N, k_1, \dots, k_s)$ denotes the number of instances I over a domain of size N that satisfy Γ and for which $|v_i(I)| = k_i$, for all $i = 1, s$.

The following is a key characterization of EM distributions.

Theorem 1. [12, page 355] Let $\Sigma = (\bar{v}, \bar{d})$ be a statistical program. For any probability distribution \bar{p} that satisfies the statistics Σ the following holds: \bar{p} is an EM distribution iff there exist parameters $\bar{\alpha}$ s.t. \bar{p} is given by the Equation (1) (equivalently: \bar{p} is associated to $(\Gamma, \bar{v}, \bar{\alpha})$).

The message of this theorem is that the weight of an instance I under the EM distribution *only depends on* $|v_i(I)|$. That is, the distribution depends exactly on the provided statistics and makes no additional assumptions. It is this property that makes the EM distribution the natural model for database statistics. In a Bayesian sense, for a fixed set of statistics the EM model yields *the optimal estimate*. We refer to Jaynes [12, page 355] for a full proof and further discussion of this point; the “only if” part of the proof is both simple and enlightening, and we include in the Appendix for completeness.

We illustrate the utility of this theorem with two simple examples:

Example 2. The Binomial-Model Consider a relation $R(A, B)$ and the statistical assertion $\#R = d$ with $\Gamma = \emptyset$. The partition function is the binomial, $T(x) = \sum_{k=0, N^2} \binom{N^2}{k} x^k = (1+x)^{N^2}$ and the EM model turns out to be the probability model that randomly inserts each tuple in R independently, with probability $p = d/N^2$. We need to check that this is an EM distribution: given an instance I of size k , $\mathbf{P}[I] = p^k (1-p)^{N^2-k}$, which we rewrite as $\mathbf{P}[I] = \omega \alpha^k$. Here $\alpha = p/(1-p)$ is the *odds* of a tuple, and $\omega = (1-p)^{N^2} = \mathbf{P}[I = \emptyset]$. This is indeed an EM distribution by Theorem 1. Asymptotic query evaluation on a generalization of this distribution to multiple tables was studied in [5]. \square

Example 3. Overlapping Ranges Consider two views⁴:

$$v_1(x, y) :- R(x, y), x < .60N \text{ and } v_2(x, y) :- R(x, y), .25N \leq x$$

and the statistical program $\#v_1 = d_1$, $\#v_2 = d_2$ (again $\Gamma = \emptyset$). Assuming $N = 100$, the views partition the domain into three buckets, $D_1 = [1, 24]$, $D_2 = [25, 59]$, $D_3 = [60, 100]$, of sizes N_1, N_2, N_3 . Here we want to say that we observe d_1 tuples in $D_1 \cup D_2$ and d_2 tuples in $D_2 \cup D_3$. The EM model gives us a precise distribution that represents only these observations and nothing more. The partition function is $(1 + x_1)^{N_1}(1 + x_1x_2)^{N_2}(1 + x_2)^{N_3}$, and the EM distribution has the form $\mathbf{P}[I] = \omega\alpha_1^{k_1}\alpha_2^{k_2}$, where $k_1 = |I \cap (D_1 \cup D_2)|$ and $k_2 = |I \cap (D_2 \cup D_3)|$; we show in Sec. 4 how to compute the parameters α_1, α_2 .

In this paper we study the *model computation problem*: given a statistical program Σ , find the parameters $\bar{\alpha}$ for the EM model. The ultimate goal of our program is to further use these parameters to estimate the size of arbitrary queries, but we will not treat the latter problem in this paper. The model depends on the size of the domain, N , and this is an undesirable property, since in practice N has no meaning other than that it is large. For that reason, we study the *asymptotic model computation problem* in this paper: find a sequence of parameters $\bar{\alpha}_N$ s.t. the distribution associated to $(\Gamma, \bar{v}, \bar{\alpha}_N)$ satisfies Σ asymptotically.

To simplify our discussion we present our results for the case when the queries in the statistical program have no range predicates, and show in Sec. 4 how to handle range predicates. Thus, from now on, until Sec. 4, we will assume all conjunctive queries to be without range predicates.

2.2 A Taxonomy for Statistical Programs

Recall that PJ denotes the class of project-join queries. We define here two subclasses. First, a *project query* is a single subgoal query without constants or repeated variables; denote P the class of project queries. Second, a *single component join query* is a project-join query with the following properties: it is minimized, has at least two subgoals, and has a single connected component; denote PJ^C the class of single component join queries. Queries V_3, V_4 in Fig. 1 are in P; queries V_5 is in PJ^C . P and PJ^C are two disjoint subclasses of PJ that do not cover PJ. Some queries in PJ are not in either class, e.g. $v(x, y) :- R(x, y), R(z, y), S(u)$ is a query that minimizes to $R(x, y), S(u)$, which is neither in P nor in PJ^C (it has two connected components): we do not treat such queries in this paper.

We classify statistical programs $\Sigma = (\bar{v}, \bar{d})$ and constraints Γ along two axes:

Definition 3. Σ is in normal form (NF) if all statistical assertions are on base tables; otherwise, it is in non-normal form (NNF).

Definition 4. (1) Σ is composite if for every statistical assertion $\#v = d$, v is either a base table or is in PJ^C . Γ is composite if for every constraint $\forall \bar{x}. w(\bar{x}) \Rightarrow R_i(\bar{x})$, w is in PJ^C . We say that Σ, Γ is composite if both are composite. (2) Σ, Γ is atomic if all their views are in P.

⁴ We represent range predicates as fractions of N so we can allow N to go to infinity.

Thus, there are four combinations of programs: NF/NNF and composite/atomic. For example, referring to Fig. 1, the program (Σ_1, Γ_1) $\Sigma_1 = \{\#R = 3000, \#T = 42000\}$ with $\Gamma_1 = \{\gamma_1\}$ is an atomic, NF program; if we add the statistical assertion (V_4, d_4) , then the program is still atomic, but no longer in normal form. On the other hand, (Σ_1, Γ) with $\Gamma = \{\gamma_3\}$ is composite and in normal form; if we add the statistic (V_5, d_5) then this becomes a composite program, not in normal form. We do not treat mixed atomic/composite programs.

3 Composite Programs

We start by discussing the case when all queries are *composite*. First, we introduce the two main techniques used in this section, conditioning on the prior, and the asymptotic probabilities from [5], then we give our results.

3.1 From Conditionals to EM Models

Recall that \mathcal{I} denotes the set of all database instances, without any constraints. Define a *prior probability distribution* to be any tuple-independent probability distribution \mathbf{P}_0 on \mathcal{I} . As seen in Example 2, this is an EM distribution for a very simple NF program, which just asserts the cardinalities of each relation, $\#R_i = d_i$, and has no constraints. Each tuple t into R_i has probability $\mathbf{P}_0[t] = d_i / N^{\text{arity}(R_i)}$, and the EM parameters are $\alpha_i = p_i / (1 - p_i) \approx p_i$. Now let's add a set of constraints Σ , i.e., consider the NF program consisting both of cardinality assertions $\#R_i$ and constraints Σ . Its EM model is obtained as follows:

Theorem 2 (Conditioning). *Let \mathbf{P} be the EM model for a NF program Σ, Γ . Then there exists a prior probability distribution \mathbf{P}_0 such that:*

$$\forall I \in \mathcal{I}(\Gamma), \quad \mathbf{P}[I] = \mathbf{P}_0[I \mid \Gamma]$$

Moreover, the expected values are obtained through the following transfer equation: $E[|q|] = E_0[|q| \mid \Gamma]$.

Proof. Let $\bar{\alpha}$ be the parameters of \mathbf{P} . Define the tuple-independent prior as follows: for each relation R_i , define $\mathbf{P}_0[t \in R_i] = p_i = \alpha_i / (1 + \alpha_i)$. (Thus, the odds of p_i are precisely α_i .) Then $\mathbf{P}_0[I] = \omega_0 \prod \alpha_i^{|R_i^I|}$ (follows by generalizing Example 2) and $\mathbf{P}[I] = \omega \prod \alpha_i^{|R_i^I|}$ (by definition). Thus, \mathbf{P} and \mathbf{P}_0 are essentially the same expression, only \mathbf{P} is defined over a restricted domain $\mathcal{I}(\Gamma) \subseteq \mathcal{I}$.

For a simple illustration, consider the statistical program $\Sigma: \#R(A, B) = d_1, \#T(B, C) = d_2$, and the constraints $\Gamma: R(x, y), R(y, z) \Rightarrow R(x, z)$ and $T(x, y), R(y, z) \Rightarrow T(z, x)$. To solve it, first solve a different, simpler program $\#R(A, B) = b_1, \#T(B, C) = b_2$, without constraints. This is a tuple-independent probability distribution \mathbf{P}_0 . Then the solution to Σ, Γ is given as $\mathbf{P}[I] = \mathbf{P}_0[I \mid \Gamma]$. The difficulty lies in choosing the statistics b_1, b_2 of the simpler model: we need to ensure that $E_0[|R| \mid \Gamma] = d_1, E_0[|T| \mid \Gamma] = d_2$.

$$\begin{aligned}
V(q) &= \text{the number of distinct variables in } q \\
a(q) &= \sum \{\text{arity}(g) \mid g \in \text{goals}(q)\} \\
D(q) &= a(q) - V(q) \\
b(q) &= \prod \{b(g) \mid g \in \text{goals}(q)\} \\
UQ(q) &= \{\eta(q) \mid \eta = \text{a substitution of variables}\} \\
E(q) &= \min D(q_0)_{q_0 \in UQ(q)} \\
UQ^0(q) &= \{q_0 \mid q_0 \in UQ(q), D(q_0) = E(q)\} \\
C(q) &= \sum_{q_0 \in UQ^0(q)} \frac{b(q_0)}{\text{aut}(q_0)}
\end{aligned}$$

Fig. 2. Notations for Theorem 3 from [5].

3.2 Background: Asymptotic Query Probabilities

Based on our discussion, we need to study prior probabilities that have the form $\mathbf{P}[t \in R] = b(R)/N^{\text{arity}(R)}$, where $b(R)$ is a constant that depends only on the relation symbol R . These tuple-independent distributions were studied in [5]. It was shown that for any Boolean conjunctive query $q \in CQ$, there exists two constants $E(q)$ and $C(q)$, which can be computed only from the constants $b(R)$ and the query expression, s.t. $\mathbf{P}[q] = C(q)/N^{E(q)} + O(1/N^{E(q)+1})$. We give the expressions for $C(q)$ and $E(q)$ in Fig. 2.

Example 4. We illustrate the notations in Fig. 2 on the query $q = R(x, y), R(y, z)$, and $b(R) = b$. $D(q) = 4 - 3 = 1$ and is called the degree of q ; and $b(q) = b^2$. $UQ(q)$ is obtained by substituting variables in q and contains four queries (up to isomorphism): q itself, then $R(x, x), R(x, y)$, then $R(x, y), R(y, y)$, and finally $R(x, x)$. Their degrees are 1, 2, 2, 1 respectively, thus $E(q) = 1$ and is called the exponent of q . $UQ^0(q)$ consists of the first and last queries (those that have $D = 1$), and $\text{aut}(q_0)$ is the number of automorphisms for q_0 , and is 1 for both queries in $UQ^0(q)$. Finally, $C(q) = b^2 + b$ is called the *coefficient* of q . Thus, $\mathbf{P}[q] = (b^2 + b)/N + O(1/N^2)$.

We consider here only conjunctive queries where all connected components have $E > 0$; this rules out some degenerate queries, whose treatment is more complex [4]. All PJ^C queries satisfy this property, since they have a single component and $E > 0$.

Theorem 3. [5] *For any conjunctive query $q \in CQ$, $\mathbf{P}_0(q) = C(q)/N^{E(q)} + O(1/N^{E(q)+1})$.*

3.3 Composite NF Programs

Theorem 4 (Composite, NF). *Consider a statistical program in normal form $\Sigma: \#R_j = d_j$, for $j = 1, m$. Consider a set of inclusion constraints Γ where all queries are composite. Then an asymptotic solution to the EM model is given by $\alpha_j = d_j/N^{\text{arity}(R_j)}$.*

The proof uses Theorem 3 and is given in the full version of the paper; it uses Theorem 3 as well as specific properties of the expressions D and E in Fig. 2. At a high level, the proof exploits the fact that $\lim_N \mathbf{P}_0[\Gamma] = 1$ (i.e., the constraints, Γ , almost surely hold), where \mathbf{P}_0 is the prior associated to the same statistical program (Σ, Γ) : that is, the constraints Γ holds almost certainly in the prior, and hence the statistics are not affected by conditioning.

Example 5. Consider the constraints $R(x, y), R(y, z) \Rightarrow R(x, z)$, and $T(x, y, z), R(y, u) \Rightarrow S(y)$, and the statistical assertions $|R| = d_1, |T| = d_2, |S| = d_3$. An asymptotic solution to the EM model is given by $\alpha_1 = d_1/N^2, \alpha_2 = d_2/N^3, \alpha_3 = d_3/N$.

3.4 Composite Non-NF Programs

Let Σ be a statistical program that consists of assertions on all relations, $\#R_j = d_j$, as well as assertions over composite views, $\#q_i = d_i$. Create a new relation symbol T_i for each statistical assertion of the same arity as the view, and define the *set equality constraints* $\forall \bar{x}. (\exists \bar{y}. q_i(\bar{x}, \bar{y}) \iff T_i(\bar{x}))$. Each set equality constraint is expressed as $\gamma_i \wedge \delta_i$, where γ_i is a full inclusion constraint and δ_i is a reverse inclusion constraint:

$$\begin{aligned}\gamma_i &\equiv \forall \bar{x}. q_i(\bar{x}) \Rightarrow T_i(\bar{x}) \\ \delta_i &\equiv \forall \bar{x}. T_i(\bar{x}) \Rightarrow (\exists \bar{y}. q_i(\bar{x}, \bar{y}))\end{aligned}$$

Denote Δ and Γ the set of all inclusion- and all reverse inclusion constraints. As before, the EM solution is given by a conditional $\mathbf{P}[I] = \mathbf{P}_0[I \mid \Delta \wedge \Gamma]$, where \mathbf{P}_0 is some tuple-independent prior. However, it is now more difficult to transfer the expected sizes, and we provide a closed form solution only for a restricted class of views.

Definition 5. *A query $q \in \mathcal{PJ}$ is a project-semi-join query if the following conditions hold. Let $\bar{x} = (x_1, \dots, x_k)$ be its head variables:*

- *q has no self-joins (i.e., no repeated relation symbol).*
- *If two different subgoals in q share a variable y , then $y \in \bar{x}$.*
- *For every subgoal g , if g contains x_i then it also contains x_{i+1} .*

A *core subgoal* is a subgoal that contains the smallest number of head variables. The core of q is the set of core subgoals and denoted G . In what follows, a transfer equation is an equation that relates a size estimate under the prior distribution to the size estimate of the distribution under constraints.

Lemma 1. *Let δ be an inverse inclusion constraint $\forall \bar{x}. T(\bar{x}) \Rightarrow \exists \bar{y}. q(\bar{x}, \bar{y})$ where q is a project-semi-join query. Define the prior:*

$$\begin{aligned}\mathbf{P}_0(t \in R_j) &= \frac{b(R_j)}{N^{\text{arity}(R_j)}} \\ \mathbf{P}_0(t \in T) &= 1 - \frac{b(T)}{N^{E(G)}}\end{aligned}$$

(Here $E(G)$ denotes the exponent of the core, see Fig. 2.) Let R_1, \dots, R_m be all subgoals in q , $m \geq 1$. Then the transfer equation for the view is:

$$E_0[|T| \mid \delta] = \frac{\prod_{R \in \text{goals}(q)} b(R)}{b(T)} \quad (2)$$

The transfer equation for any other relation in R is as follows. If the query consists only of the core, then:

$$E_0[|R_i| \mid \delta] = b(R_i) + \frac{C(G)}{b(T)} \quad (3)$$

($C(G)$ is the coefficient of G , see Fig. 2.) If the query has subgoals other than the core, then the expected cardinalities are unchanged.

We prove the lemma in the full version of the paper. From the lemma we derive:

Theorem 5 (Composite, non-NF). *Let Σ be a statistical program where all queries are project-semi-join queries, and do not share common subgoals. Then an EM model for Σ has the following parameters:*

- For every base relation R_j , the parameter is $\alpha_j = b_j / N^{\text{arity}(R_j)}$.
- For every view assertion v_j , the parameter is $\alpha_j = N^{E(G)} / b_j$, where G is the core of v_j .

where the numerical values b_j are obtained by solving a system of equations (2) and (3).

It is interesting to compare the solution to an NF program to that of a non-NF program (Theorems 4 and 5). For NF programs all parameters have the form $\alpha_i = d / N^a$, for integer $a > 0$. For non-NF programs some parameters have the form N^a / d and, thus, go to infinity.

Example 6. Consider the following statistical program⁵:

$$\begin{aligned} \#R_1 &= d_1 \quad \#R_2 = d_2 \\ v(x) &:- R_1(x), R_2(x) \quad \#v = d_3 \end{aligned}$$

Thus, we are given the sizes of R_1 , R_2 , and of their intersection. We introduce a new relation symbol T and the constraint $\delta = T(x) \Leftrightarrow R_1(x), R_2(x)$, then define the program in normal form:

$$\#R_1 = d_1 \quad \#R_2 = d_3 \quad \#T = d_3$$

The theorem gives us the EM solution as follows. The core is the entire query, hence we define the prior:

$$\mathbf{P}_0(R_1(a)) = e_1 / N \quad \mathbf{P}_0(R_2(a)) = e_2 / N \quad \mathbf{P}_0(T(a)) = 1 - e_3 / N$$

⁵ Its partition function is $(1 + x_1 + x_2 + x_1x_2)^N$. Intuitively, this is because each of the n tuples is in R_1 and so pays x_1 , is in R_2 and so pays x_2 or is in both R_1 and R_2 and so pays x_1x_2 .

where $\mathbf{P}_0(R_1(a))$ denotes the marginal probability of the tuple $R_1(a)$. Note that T has a very large probability. This gives us an EM model to our initial statistical program if we solve e_1, e_2, e_3 in:

$$\begin{aligned} d_3 &= \frac{e_1 e_2}{e_3} \\ d_1 &= e_1 + \frac{e_1 e_2}{e_3} \\ d_2 &= e_2 + \frac{e_1 e_2}{e_3} \end{aligned}$$

Example 7. A more complex example is a statistical program that uses the following project-semi-join view:

$$\begin{aligned} v_6(x_1, x_2, x_3) \quad :- \quad & R_1(x_1, x_2, x_3, y), R_2(x_2, x_3), R_3(x_2, x_3, z), \\ & R_4(x_1, x_2, x_3), R_5(x_1, x_2, x_3) \end{aligned}$$

The core consists of R_2 and R_3 , and so we define $\mathbf{P}_0[t \in T] = e_T/N^2$, where e_T is chosen such that $d_2 d_3 / e_T = d_6$.

4 Bucketization

Finally, we re-introduce range predicates like $x < c$, both in the constraints and in the statistical assertions. To extend the asymptotic analysis, we assume that all constants are expressed as fractions of the domain size N , e.g., in Ex. 3 we have $v_1(x, y) :- R(x, y), x < 0.25N$.

Let $\bar{R} = R_1, \dots, R_m$ be a relational schema, and consider a statistical program Σ, Γ with range queries, over the schema \bar{R} . We translate it into a *bucketized* statistical program Σ^0, Γ^0 , over a new schema \bar{R}^0 , as follows. First, use all the constants that occur in the constraints or in the statistical assertions to partition the domain into b buckets, $D = D_1 \cup D_2 \cup \dots \cup D_b$. Then define as follows:

- For each relation name R_j of arity a define b^a new relation symbols, $R_j^{i_1 \dots i_a} = R_j^{\bar{i}}$, where $i_1, \dots, i_a \in [b]$; then \bar{R}^0 is the schema consisting of all relation names $R_j^{i_1 \dots i_a}$.
- For each conjunctive query q with range predicates, denote $\mathbf{buckets}(q) = \{q^{\bar{i}} \mid \bar{i} \in [b]^{|Vars(q)|}\}$ the set of queries obtained by associating each variable in q to a unique bucket, and annotating the relations accordingly. Each query in $\mathbf{buckets}(q)$ is a conjunctive query over the schema \bar{R}^0 , without range predicates, and q is logically equivalent to their union.
- Let $BV = \bigcup \{\mathbf{buckets}(v) \mid (v, d) \in \Sigma\}$ (we include in BV queries up to logical equivalence), and let c_u denote a constant for each $u \in BV$, s.t. for each statistical assertion $\#v = d$ in Σ the following holds

$$\sum_{u \in \mathbf{buckets}(v)} c_u = d \tag{4}$$

Denote Σ^0 the set of statistical assertions $\#u = c_u, u \in BV$.

- For each inclusion constraint $w \Rightarrow R$ in Γ , create $b^{|Vars(w)|}$ new inclusion constraints, of the form $w^{\bar{j}} \Rightarrow R^{\bar{i}}$; call Γ^0 the set of new inclusion constraints.

Then the following holds:

Proposition 1. *Let Σ^0, Γ^0 be the bucketized program for Σ, Γ . Let $\bar{\beta} = (\beta_k)$ be the EM model of the bucketized program. Consider some parameters $\bar{\alpha} = (\alpha_j)$. Suppose that for every statistical assertion $\#v_j = d_j$ in Σ condition (4) holds, and the following condition holds for every query $u_k \in BV$:*

$$\beta_k = \prod_{j: u_k \in buckets(v_j)} \alpha_j \quad (5)$$

Then $\bar{\alpha}$ is a solution to the EM model for Σ, Γ .

This gives us a general procedure for solving the EM model for programs with range predicates: introduce new unknowns c_j^i and add Equations (4) and (5), then solve the EM model for the bucketized program under these new constraints.

Example 8. Recall Example 3: we have two statistics $\#\sigma_{A \leq 0.60N}(R) = d_1$, and $\#\sigma_{A \geq 0.25N}(R) = d_2$. The domain D is partitioned into three domains, $D_1 = [1, 0.25N)$, $D_2 = [0.25N, 0.60N)$, and $D_3 = [0.60N, N]$, and we denote N_1, N_2, N_3 their sizes. The bucketization procedure is this. Define a new schema R^1, R^2, R^3 , with the statistics $\#R^1 = c^1$, $\#R^2 = c^2$, $\#R^3 = c^3$, then solve it, subject to the Equations (5):

$$\begin{aligned} \beta_1 &= \alpha_1 \\ \beta_2 &= \alpha_1 \alpha_2 \\ \beta_3 &= \alpha_2 \end{aligned}$$

We can solve for R^1, R^2, R^3 , since each R^i is given by a binomial distribution with tuple probability $\beta_i / (1 + \beta_i) = c^i / N_i$. Now use Equations (4), $c^1 + c^2 = d_1$ and $c^2 + c^3 = d_2$ to obtain:

$$\begin{aligned} N_1 \frac{\alpha_1}{1 + \alpha_1} + N_2 \frac{\alpha_1 \alpha_2}{1 + \alpha_1 \alpha_2} &= d_1 \\ N_3 \frac{\alpha_2}{1 + \alpha_2} + N_2 \frac{\alpha_1 \alpha_2}{1 + \alpha_1 \alpha_2} &= d_2 \end{aligned}$$

Solving this gives us the EM model. Consistent histograms [16] had a similar goal of using EM to capture statistics on overlapping intervals, but use a different, simpler probabilistic model based on frequencies.

5 Related Work

There are two bodies of work that are most closely related to this paper. The first consists of the work in cardinality estimation. As noted above, while a variety of synopsis structures have been proposed for cardinality estimation [1, 8, 10, 15], they have all focused on various sub-classes of queries and deriving estimates for

arbitrary query expressions has involved *ad-hoc* steps such as the independence and containment assumptions which result in large estimation errors [11]. In contrast, we ask the question what is the framework for performing cardinality estimation over arbitrary expressions in the presence of incomplete information. We approach this task via the EM principle.

The EM model has been applied in prior work to the problem of cardinality estimation [14, 16]. However, the focus was restricted to queries that consist of conjunctive selection predicates over single tables. In contrast, we explore a full-fledged EM model that can incorporate statistics involving arbitrary first-order expressions.

Another body of related work consists of the work in probabilistic databases [7] which focuses on efficient query evaluation over a probabilistic database. The input statistics impose many possible distributions over the *possible worlds* and we choose the distribution that has maximum entropy. Our focus in this paper is in deriving the parameters of this EM distribution. The related problem of query estimation for a given model is not addressed in this paper. This is closely related to the problem of evaluating queries over probabilistic databases.

Finally, we observe that entropy-maximization is a well-established principle in statistics for handling incomplete information [12]. As with probabilistic databases, new challenges emerge in the context of database systems, in our case the nature of statistics.

6 Conclusion

In this paper we propose to model arbitrary database statistics using an Entropy-Maximization probability distribution. This model is attractive because any query has a well-defined size estimate, all statistics are treated as a whole rather than as individual synopses, and the model extends smoothly when new statistics are added. We reported in this paper several results that give explicit asymptotic solutions to statistical programs in several cases. As part of our technical development we described a technique encapsulated as the conditioning theorem (Theorem 2) that is of independent interest and are likely to be applicable to other statistical programs.

We are leaving for future work the second part: using an EM model to obtain query size estimates. This has been solved in the past only for the independent case [6].

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A Proof of Theorem 1

The “only if” direction is very simple to derive by using the Lagrange multipliers for solving:

$$F_0 = \sum_{I \in \mathcal{I}} p_I - 1 = 0 \quad (6)$$

$$\forall i = 1, \dots, s : F_i = \sum_{I \in \mathcal{I}} |v_i(I)| p_I - d_i = 0 \quad (7)$$

$$H = \text{maximum, where } H = \sum_{I \in \mathcal{I}} p_I \log p_I \quad (8)$$

According to that method, one has to introduce $s + 1$ additional unknowns, $\lambda, \lambda_1, \dots, \lambda_s$: an EM distribution is a solution to a system of $|\mathcal{I}| + s + 1$ equations consisting of Eq.(6), (7), and the following $|\mathcal{I}|$ equations:

$$\forall I \in \mathcal{I} : \frac{\partial(H - \sum_{i=0,s} \lambda_i G_i)}{\partial p_I} = \log p_I - (\lambda_0 + \sum_{i=1,s} \lambda_i |v_i(I)|) = 0$$

This implies $p_I = \exp(\lambda_0 + \sum_{i=1,s} \lambda_i |v_i(I)|)$, and the claim follows by denoting $\omega = \exp(\lambda_0)$, and $\alpha_i = \exp(\lambda_i)$, $i = 1, s$.