

# Moment-Based Quantile Sketches for Efficient High Cardinality Aggregation Queries

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## ABSTRACT

Interactive analytics increasingly involves querying for quantiles over sub-populations of high cardinality datasets. Data processing engines such as Druid and Spark use mergeable summaries to estimate quantiles, but summary merge times can be a bottleneck during aggregation. We show how a compact and efficiently mergeable quantile sketch can support aggregation workloads. This data structure, which we refer to as the moments sketch, operates with a small memory footprint (200 bytes) and computationally efficient (50ns) merges by tracking only a set of summary statistics, notably the sample moments. We demonstrate how we can efficiently estimate quantiles using the method of moments and the maximum entropy principle, and show how the use of a cascade further improves query time for threshold predicates. Empirical evaluation shows that the moments sketch can achieve less than 1 percent quantile error with  $15\times$  less overhead than comparable summaries, improving end query time in the MacroBase engine by up to  $7\times$  and the Druid engine by up to  $60\times$ .

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## 1. INTRODUCTION

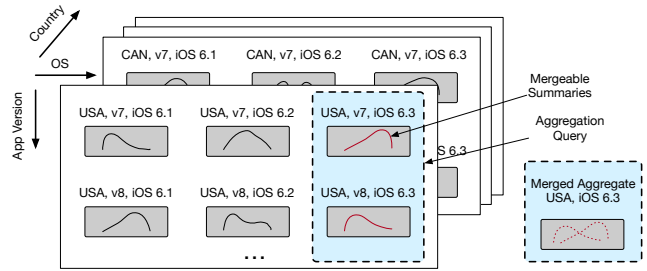
Performing interactive multi-dimensional analytics over data from sensors, devices, and servers increasingly requires computing aggregate statistics for specific subpopulations and time windows [4, 29, 65]. In applications such as A/B testing [38, 42], exploratory data analysis [8, 74], and operations monitoring [2, 14], analysts perform aggregation queries to understand how specific user cohorts, device types, and feature flags are behaving. In particular, computing quantiles over these subpopulations is an essential part of debugging and real-time monitoring workflows [26].

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**Figure 1:** Given a data cube with pre-aggregated summaries, we can compute roll-ups along specific dimensions by merging the relevant summaries. Efficiently mergeable summaries enable scalable aggregations.

As an example of this quantile-driven analysis, our collaborators on a Microsoft application monitoring team collect billions of telemetry events daily from millions of heterogeneous mobile devices. Each device tracks multiple metrics including request latency and memory usage, and is associated with dimensional metadata such as application version and hardware model. Engineers issue quantile queries on a Druid-like [82] in-memory data store, aggregating across different dimensions to monitor their application (e.g., examine memory trends across device types) and debug regressions (e.g., examine tail latencies across versions). Querying for a single percentile in this deployment can require aggregating hundreds of thousands of dimension value combinations.

When users wish to examine the quantiles of specific slices of a dataset, OLAP engines such as Druid and Spark support computing approximate quantiles using compressed representations (*summaries*) of the data values [39, 67, 82, 83]. By pre-aggregating a summary for each combination of dimension values, Druid and similar engines can reduce query times and memory usage by operating over the relevant summaries directly, effectively constructing a data cube [33, 65].

Given a time interval and a metric with  $d$  associated dimensions, Druid maintains one pre-aggregated summary for each  $d$ -tuple of dimension values. These summaries are kept in RAM across a number of nodes, with each node scanning relevant summaries to process subsets of the data specified by a user query. Figure 1 illustrates how these *mergeable* [3] summaries can be aggregated to compute quantile roll-ups along different dimensions without scanning over the raw data.

More concretely, a Druid-like data cube in our Microsoft deployment with 6 dimension columns, each with 10 distinct

values, is stored as a set of up to  $10^6$  summaries per time interval. On this cube, computing the 99-th percentile latency for a specific app version can require 100,000 merges, or even more for aggregation across complex time ranges. When there are a limited number of dimensions but enormous data volumes, it is cheaper to maintain these summaries than scan over billions of raw datapoints.

Many quantile summaries support the merge operation [3, 28, 34], but their runtime overheads can lead to severe performance penalties on high-cardinality datasets. Based on our experiments (Section 6.2.1), one million 1KB GK-sketches [34] require more than 3 seconds to merge sequentially, limiting the types of queries users can ask interactively. The merging can be parallelized, but additional worker threads still incur coordination and resource usage overheads. Materialized views [37, 44, 51, 61], sliding window sketches [25], and dyadic intervals can also reduce this overhead. However, dyadic intervals only apply to ordered dimensions and maintaining materialized views for multiple dimension roll-ups can be prohibitively expensive in a real-time stream, so merge time remains a relevant bottleneck.

In this paper, we enable interactive quantile queries over high-cardinality aggregates by introducing a compact and efficiently mergeable quantile sketch and associated quantile estimation routines. We draw a connection between the classic *method of moments* for parameter estimation in statistics [79] and the need for efficient summary data structures. We show that storing the sample moments  $\mu_i = \frac{1}{n} \sum x^i$  and log-moments  $\nu_i = \frac{1}{n} \sum \log^i(x)$  can enable accurate quantile estimation over a range of real-world datasets while utilizing fewer than 200 bytes of memory and incurring merge times of less than 50 nanoseconds. In the context of quantile estimation, we refer to our proposed summary data structure as the *moments sketch*.

While constructing the moments sketch is straightforward, the inverse problem of estimating quantiles from the summary is more complex. The statistics in a moments sketch provide only loose constraints on the distribution of values in the original dataset: many distributions might match the moments of a moments sketch but fail to capture the dataset structure. Therefore, we make use of the *principle of maximum entropy* [41] to compute a “least-biased” quantile estimate for a moments sketch. On continuous real-valued datasets, we find that this approach yields more accurate estimates than alternative methods, achieving  $\epsilon \leq 1\%$  error with 200 bytes of memory. To achieve this, we also describe a series of practical optimizations to standard entropy maximization that allow us to compute quantile estimates in under 1 millisecond on a range of real-world datasets.

These query times make the moments sketch a suitable summary when many merges (hundreds of thousands) are required, memory per-summary may be limited to less than 1 kilobyte, and  $\epsilon = .01$  error is acceptable. The moments sketch and our maximum entropy estimate is most useful in datasets without strong discretization and when very small  $< 10^{-4}$  error is not required. The maximum entropy principle is less accurate when there are clusters of discrete values in a dataset (Section 6.2.3), and floating point stability (Section 4.3.2) limits the minimum achievable error using this approach.

Moving beyond simple quantile queries, many complex queries depend on the quantile estimates of multiple subpopulations. For example, data exploration systems such

as MacroBase [8] are interested in finding all subpopulations that match a given threshold condition (e.g., subpopulations where the 95th percentile latency is greater than the global 99th percentile latency). Given a large number of subpopulations, the cost of millisecond-level quantile estimates on thousands of subgroups will accumulate. Therefore, to support threshold queries over multiple populations, we extend our quantile estimator with a *cascade* [75], or sequence of increasingly precise and increasingly expensive estimates based on bounds such as the Markov inequalities. For queries with threshold conditions, the cascades dramatically reduce the overhead of quantile estimation in a moments sketch, by up to  $25\times$ .

We implement the moments sketch both as a reusable library and as part of the Druid and MacroBase analytics engines. We empirically compare its accuracy and efficiency with alternative mergeable quantile summaries on a variety of real-world datasets. We find that the moments sketch offers 16 to  $50\times$  faster merge times than alternative summaries with comparable accuracy. This enables 15 to  $50\times$  faster query times on real datasets. Moreover, the moments sketch enables up to  $7\times$  faster analytics queries when integrated with MacroBase and  $60\times$  faster end-to-end queries when integrated with Druid.

In summary, we make the following contributions:

- We illustrate how statistical moments are useful as efficient mergeable quantile sketches in aggregation and threshold-based queries over high-cardinality data.
- We demonstrate how statistical and numerical techniques allow us to solve for accurate quantile estimates in less than 1 ms, and show how the use of a cascade further improves estimation time on threshold queries by up to  $25\times$ .
- We evaluate the use of moments as quantile summaries on a variety of real-world datasets and show that the moments sketch enables 15 to  $50\times$  faster query times in isolation, up to  $7\times$  faster queries when integrated with MacroBase and up to  $60\times$  faster queries when integrated with Druid over comparably-accurate quantile summaries.

The remainder of this paper proceeds as follows. In Section 2, we discuss related work. In Section 3, we review relevant background material. In Section 4, we describe the proposed moments sketch. In Section 5, we describe a cascade-based approach for efficiently answering threshold-based queries. In Section 6, we evaluate the moments sketch in a series of microbenchmarks. In Section 7, we evaluate the moments sketch as part of the Druid and MacroBase systems, and benchmark its performance in a sliding window workflow. We conclude in Section 8. We include supplemental appendices in an extended technical report [30].

## 2. RELATED WORK

**High-performance aggregation.** The aggregation scenarios in Section 1 are found in many existing streaming data systems [8, 16, 24, 65, 82], as well as data cube [33, 69], data exploration [2], and visualization [17] systems. In particular, these systems are can perform interactive aggregations over time windows and along many cube dimensions, motivating the design of our sketch. Many of these systems

use approximate query processing, sampling, and summaries to improve query performance [4, 35, 59], but do not develop data structures specific to quantiles. We believe the moments sketch serves as a useful primitive in these engines.

Sensor networking is a rich source of algorithms for heavily resource-constrained settings. Sensor network aggregation systems [53] support large scale roll-ups, but work in this area is largely focused on the complementary problem of communication plans over a network [21, 45, 54]. Mean, min, max, and standard deviation in particular are used in [53] as functions amenable to computation-constrained environments, but the authors do not consider higher moments or their application to quantile estimation.

Several database systems make use of summary statistics in general-purpose analytics. Muthukrishnan et al [60] observe that the moments are a convenient statistic in streaming settings and use it to fill in missing integers. Data Canopy [78] uses first and second moments as an efficiently mergeable statistic for computing standard deviations and linear regressions. Similarly, systems on probabilistic data cubes such as [81] use the first and second moments to prune queries over cube cells that store distributions of values. In addition, many methods use compressed data representations to perform statistical analyses such as linear regression, logistic regression, and PCA [19, 63, 70, 80]. We are not aware of prior work utilizing higher moments to efficiently estimate quantiles for high-dimensional aggregation queries.

**Quantile summaries.** There are a variety of summary data structures for the  $\epsilon$ -approximate quantile estimation problem [18, 23, 34, 71]. Some of these summaries assume values from a fixed universe [23, 71], while others operate using only comparisons [3, 34]. Our proposed moments sketch and others [12, 28] operate on real values. Agarwal et al. [3] provide the initial motivation for mergeable summaries, as well as a proposed mergeable quantile sketch. The authors in [52, 77] benchmark a variety of quantile summaries but do not directly evaluate merge time. Zhuang [84] evaluates merge performance of a variety of quantile summaries in a distributed setting, finding the **Random** summary to be the fastest. To our knowledge we are the first to introduce and evaluate the moments sketch for fast merge times and low space overhead.

**Method of moments.** The method of moments is a well-established statistical technique for estimating the parameters of probability distributions [79]. The main idea behind this approach is that the parameters of a distribution of interest  $P$  can be related to the expectations of functions of the random variable  $X \sim P$ . As a general method for consistent statistical parameter estimation, the method of moments is used across a wide range of fields, including econometrics [36], physics [32, 57], and machine learning [7, 11, 43]. In this work, we demonstrate how the method of moments, applied in conjunction with practical performance optimizations, can scale to support real-world latency-sensitive query processing workloads.

**Maximum entropy principle.** The maximum entropy principle prescribes that one should select the least informative distribution that is consistent with the observed data. In the database community, this principle has been applied to estimating cardinality [73] and predicate selectivity [55]. Mead and Papanicolaou [57] apply the maximum entropy principle to the problem of estimating distributions subject

to moment constraints; follow-up work proposes the use of Chebyshev polynomials for stability [10, 72] and faster approximation algorithms [9], though we have not seen any practical implementations suitable for use in a database. The maximum entropy principle is also used in machine learning, notably in the context of *maximum entropy models* [13]. For example, in natural language processing, maximum entropy models are a popular choice for tasks such as text classification [62] and sequence tagging [48].

### 3. BACKGROUND

In this section, we review the approximate quantile estimation problem, mergeable quantile summaries, and our target query cost model.

#### 3.1 Quantile Queries

Given a dataset  $D$  with  $n$  elements, for any  $\phi \in (0, 1)$ , the  $\phi$ -quantile of  $D$  is the item  $x \in D$  with rank  $r(x) = \lfloor \phi n \rfloor$ , where the rank of an element  $x$  is the number of elements in  $D$  smaller than  $x$ .

An  $\epsilon$ -approximate  $\phi$ -quantile is an element with rank between  $(\phi - \epsilon)n$  and  $(\phi + \epsilon)n$  [3]. Given an estimated  $\phi$ -quantile  $\hat{q}_\phi$ , we can also define its *quantile error*  $\epsilon$  [52] as the following:

$$\epsilon = \frac{1}{n} |\text{rank}(\hat{q}_\phi) - \lfloor \phi n \rfloor|, \quad (1)$$

such that an  $\epsilon$ -approximate quantile has error at most  $\epsilon$ . For example, given a dataset  $D = \{1, \dots, 1000\}$ , an estimate  $\hat{q}_{0.5} = 504$  for the  $\phi = 0.5$  quantile would have error  $\epsilon = 0.003$ . In this paper, we consider datasets  $D$  represented by collections of real numbers  $D \subset \mathbb{R}$ .

*Quantile summaries* are data structures that provide approximate quantile estimates for a dataset given space sub-linear in  $n$ . These summaries usually have a parameter  $k_\epsilon$  that trades off between the size of the summary and the accuracy of its estimates. An  $\epsilon$ -approximate quantile summary provides  $\epsilon$  approximate  $\phi$ -quantiles, where  $\epsilon$  can be a function of space usage and the dataset [18, 23, 34, 71].

#### 3.2 Mergeable Summaries

Agarwal et al. [3] introduce the concept of *mergeability* to accurately combine summaries in distributed settings. Formally, for a summary with parameter  $k_\epsilon$ , we use  $S(D, k_\epsilon)$  to denote a valid summary for a dataset  $D$ . For any pair of datasets  $D_1$  and  $D_2$ , the summarization routine  $S$  is *mergeable* if there exists an algorithm (i.e., the “merge” procedure) that produces a combined summary

$$S(D_1 \uplus D_2, k_\epsilon) = \text{merge}(S(D_1, k_\epsilon), S(D_2, k_\epsilon))$$

from any two input summaries, where  $\uplus$  denotes multiset addition.

Intuitively, a summary is mergeable if there is no accuracy cost to combining pre-computed summaries compared with constructing a summary on the raw data. Thus, mergeable summaries are *algebraic* aggregate functions in the data cube literature [33]. As an example, an equi-depth histogram [22] on its own is not mergeable because there is no way to accurately combine two overlapping histogram buckets without access to additional data.

Mergeable summaries can be incorporated naturally into a variety of distributed systems. In the MapReduce paradigm, a “map” function can construct summaries over shards while

a “reduce” function merges them to summarize a complete dataset [3]. In the GLADE system [68], mergeable summaries are an example of a *Generalized Linear Aggregate* (GLA), a user-defined computation that can be incrementally aggregated across nodes.

### 3.3 Query Model

As described in Section 1, we focus on improving the performance of quantile queries over aggregations on high cardinality datasets. Given a dataset with  $d$  categorical dimensions, we consider data cubes that maintain summaries for every  $d$ -way dimension value tuple as one natural setting for high performance aggregations, and many other settings are also applicable [78]. In these settings, query time is heavily dependent on the number of merges and the time per merge.

We consider two broad classes of queries in this paper. First, *single quantile* queries ask for quantile estimates for a single specified population. For example, we can query the 99th percentile of latency over the last two weeks for a given version of the application:

```
SELECT percentile(latency, 99) FROM requests
WHERE time > date_sub(curdate(), 2 WEEK)
AND app_version = "v8.2"
```

To process this query in time  $t_{\text{query}}$ , we would need to merge  $n_{\text{merge}}$  summaries, each with runtime overhead  $t_{\text{merge}}$ , and then estimate the quantile from the merged summary with runtime cost  $t_{\text{est}}$ . This results in total query time:

$$t_{\text{query}} = t_{\text{merge}} \cdot n_{\text{merge}} + t_{\text{est}}. \quad (2)$$

We evaluate the different regimes where queries are bottlenecked on merges and estimation in Figure 6 in Section 6.2.2: merge time begins to dominate at around  $n_{\text{merge}} \geq 10^4$ .

We also consider *threshold* queries which are conditioned on sub-groups or windows with percentiles above a specified threshold. For example, we may be interested in combinations of application version and hardware platform for which the 99th percentile latency exceeds 100ms:

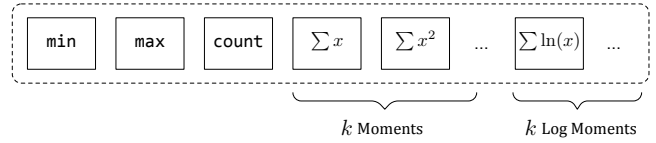
```
SELECT app_version, hw_model,
       PERCENTILE(latency, 99) as p99
FROM requests
GROUP BY app_version, hw_model
HAVING p99 > 100
```

Such queries are very useful for debugging and data exploration [8], but have additional runtime cost that depends on the number of groups  $n_{\text{groups}}$  since  $t_{\text{est}}$  can be significant when one is searching for high quantiles over thousands of sub-groups. This results in total query time:

$$t_{\text{query}} = t_{\text{merge}} \cdot n_{\text{merge}} + t_{\text{est}} \cdot n_{\text{groups}}. \quad (3)$$

## 4. THE MOMENTS SKETCH

In this section, we describe how we perform quantile estimation using the moments sketch. First, we review the summary statistics stored in the moments sketch and describe how they comprise an efficiently mergeable sketch. Second, we describe how we can use the method of moments and the maximum entropy principle to estimate quantiles from the moments sketch, with details on how we resolve practical difficulties with numerical stability and estimation time. We conclude with a discussion of theoretical guarantees on the approximation error of quantiles estimated from the sketch.



**Figure 2:** The moments sketch is an array of floating point values.

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#### Algorithm 1: Moments sketch operations

---

```
input: number of moments  $k$ 
function Init( $x$ )
     $x_{\min}, x_{\max} \leftarrow \infty, -\infty$ 
     $n, \vec{\mu}, \vec{\nu} \leftarrow 0, \vec{0}$ 
function Accumulate( $x$ )
     $x_{\min}, x_{\max} \leftarrow \min\{x, x_{\min}\}, \max\{x, x_{\max}\}$ 
     $n \leftarrow n + 1$ 
    for  $i \in \{1, \dots, k\}$  do
         $\mu_i \leftarrow \frac{n-1}{n}\mu_i + \frac{1}{n}x^i$   $\triangleright$  Standard moments
        if  $x > 0$  then
             $\nu_i \leftarrow \frac{n-1}{n}\nu_i + \frac{1}{n}\log^i(x)$   $\triangleright$  Log-moments
function Merge( $o$ )  $\triangleright o$  another sketch
     $x_{\min} \leftarrow \min\{o.x_{\min}, x_{\min}\}$ 
     $x_{\max} \leftarrow \max\{o.x_{\max}, x_{\max}\}$ 
     $n, \vec{\mu}, \vec{\nu} \leftarrow n + o.n, \vec{\mu} + o.\vec{\mu}, \vec{\nu} + o.\vec{\nu}$ 
```

---

### 4.1 Moments Sketch Statistics

The moments sketch of a dataset  $D$  is a set of floating point values: the minimum value  $x_{\min}$ , the maximum value  $x_{\max}$ , the count  $n$ , the sample moments  $\mu_i = \frac{1}{n} \sum_{x \in D} x^i$  and the sample logarithmic moments  $\nu_i = \frac{1}{n} \sum_{x \in D} \log^i(x)$  for  $i \in \{1, \dots, k\}$  (Figure 2). The moments sketch has an integer parameter  $k$ , which describes the highest power used in the moments. We refer to  $k$  as the *order* of a moments sketch. Each sample moment provides additional information about the distribution, so higher-order moments sketches are more precise but have higher space and computation time overheads.

The moments sketch supports a number of basic operations: **init** creates an empty sketch, **accumulate** updates the sketch via point-wise additions, and **merge** updates the sketch by merging it with another moments sketch. One can construct a moments sketch either by using **accumulate** to incrementally add raw data points, or using **merge** on preaggregated existing sketches. When accumulating elements point-wise, we update the minimum and maximum, then add to the counts and moments. As an implementation detail, we can accumulate the unscaled sums  $\sum x^i$  and  $\sum \log^i(x)$  internally instead of the  $\mu_i, \nu_i$ . We merge two moments sketches by combining the minimum, maximum, count, and the moments via comparison and potentially vectorized addition. This merge operation preserves the property that a moments sketch constructed using only **accumulate** is identical (up to floating point precision) to a moments sketch constructed from merging existing sketches of partitions of the data, so there is no accuracy loss in pre-aggregating. We provide pseudocode for these in Algorithm 1. The moments sketch additionally supports quantile estimation routines described in Section 4.2 in order to answer end-user queries.

The moments sketch thus supports all of the basic user-defined aggregate operations [20, 68].

**Log-moments.** The moments sketch records logarithmic moments (log-moments) in order to recover better quantile estimates for long-tailed datasets. In particular, taking the logarithm of data points is useful when values in the dataset can vary over several orders of magnitude. In general, when updating a moments sketch in a streaming manner or when maintaining multiple moments sketches in a distributed setting, we cannot know *a priori* whether standard moments or log-moments are more appropriate for the given dataset. Therefore, our default approach is to store both sets of moments up to the same order  $k$ . Given additional prior knowledge of the data distribution, we may choose to maintain a moments sketch with only a single set of moments.

Data points with negative values pose a potential problem for the log-moments since the logarithm is undefined for these points. There are several strategies for addressing this, including storing separate sums for positive and negative values and shifting all values so that they are positive. In this paper, we adopt the simple approach of ignoring the log sums when there are any negative values, and computing estimates using the remaining statistics.

**Remark on pathological distributions.** The moments of certain “pathological” distributions may be undefined; for example, the Cauchy distribution  $f(x) = \pi^{-1}(1+x^2)^{-1}$  does not have finite moments of any order. However, the moments sketch tracks the moments of an empirical dataset, which are always well-defined. This suits our goal of estimating quantiles of a finite dataset.

## 4.2 Estimating Quantiles

**Method of moments.** To estimate quantiles from a moments sketch, we apply the *method of moments* [7, 11, 43, 79] to construct a distribution  $f(x)$  whose moments match those recorded in the sketch. Specifically, given a moments sketch with minimum  $x_{\min}$  and maximum  $x_{\max}$ , we solve for a pdf  $f(x)$  supported on  $[x_{\min}, x_{\max}]$  with moments equal to the values in the moments sketch.

$$\int_{x_{\min}}^{x_{\max}} x^i f(x) dx = \mu_i \quad \int_{x_{\min}}^{x_{\max}} \log^i(x) f(x) dx = \nu_i$$

We can then report the quantiles of  $f(x)$  as estimates for the quantiles of the dataset.

In general, a finite set of moments does not uniquely determine a distribution [5]. That is, there are often many possible distributions with varying quantiles that each match a given set of sample moments. Therefore, we must disambiguate between them.

**Maximum entropy.** In this work, we use the *principle of maximum entropy* [41] to select a unique distribution that satisfies the given moment constraints. Intuitively, the differential Shannon entropy  $H$  of a distribution with pdf  $f(x)$ , defined as  $H[f] = -\int_{\mathcal{X}} f(x) \log f(x) dx$ , is a measure of the degree of *uninformativeness* of the distribution. For example, a uniform distribution has higher entropy than a point mass distribution. Thus, the maximum entropy distribution can be seen as the distribution that encodes the *least* additional information about the data beyond that captured by the given moment constraints.

Applying the maximum entropy principle to the moments

sketch, we estimate quantiles by solving for the pdf  $f$  that maximizes the entropy while matching the moments in the sketch. Following, we estimate quantiles using numeric integration and the Brent’s method for root finding [64].

In practice, we find that the use of maximum entropy distributions yields quantile estimates with comparable accuracy to alternative methods on a range of real-world datasets, unless the datasets are more discrete than continuous. We discuss our empirical results further in Section 6.2.3.

**Optimization.** We now describe how to solve for the maximum entropy distribution  $f$ . We trade off between accuracy and estimation time by solving for  $f$  subject to the first  $k_1$  standard moments and  $k_2$  log-moments stored in the sketch; incorporating more moments leads to more precise estimates but more computationally expensive estimation. As previously noted, for datasets with non-positive values (i.e.,  $x_{\min} \leq 0$ ), we set  $k_2 = 0$ . Therefore, our goal is to find the solution  $f$  to the following optimization problem:

$$\begin{aligned} & \underset{f \in \mathcal{F}[x_{\min}, x_{\max}]}{\text{maximize}} && H[f] && (4) \\ & \text{subject to} && \int_{x_{\min}}^{x_{\max}} x^i f(x) dx = \mu_i, && i \in \{1, \dots, k_1\} \\ & && \int_{x_{\min}}^{x_{\max}} \log^i(x) f(x) dx = \nu_i, && i \in \{1, \dots, k_2\} \end{aligned}$$

where  $\mathcal{F}[x_{\min}, x_{\max}]$  denotes the set of distributions supported on  $[x_{\min}, x_{\max}]$ .

It is well known that the solution to Problem (4) is a member of the class of exponential family distributions [41]:

$$f(x; \theta) = \exp \left( \theta_0 + \sum_{i=1}^{k_1} \theta_i x^i + \sum_{i=1}^{k_2} \theta_{k_1+i} \log^i(x) \right),$$

where  $\theta_0$  is a normalization constant such that  $f(x; \theta)$  integrates to 1 over the domain  $[x_{\min}, x_{\max}]$ . The maximum entropy distribution is determined by the parameter  $\theta$  such that  $f(x; \theta)$  satisfies the moment constraints in Problem (4).

In order to solve for  $\theta$ , we define the potential function  $L(\theta)$  from [57]:

$$\begin{aligned} L(\theta) = & \int_{x_{\min}}^{x_{\max}} \exp \left( \sum_{i=0}^{k_1} \theta_i x^i + \sum_{i=1}^{k_2} \theta_{k_1+i} \log^i x \right) dx && (5) \\ & - \theta_0 - \sum_{i=0}^{k_1} \theta_i \mu_i - \sum_{i=1}^{k_2} \theta_{k_1+i} \nu_i \end{aligned}$$

$L(\theta)$  is a convex function over  $\theta$  and is constructed so that the minimizing solution  $\theta_{\text{opt}} = \arg \min_{\theta \in \mathbb{R}^{k_1+k_2-1}} L(\theta)$  is exactly the set of coefficients which satisfy the constraints in Problem (4). Equation (5) thus transforms the constrained optimization problem in (4) into an unconstrained convex optimization problem which we solve using Newton’s method [15]. Formulae for the gradient and Hessian of Equation (5) are provided in Appendix A in [30].

Instead of Newton’s method, one could also use first-order optimization routines such as SGD and BFGS [50] to solve Problem (4): these would not use the Hessian but might require more steps to achieve convergence. As we describe in Section 4.3, each additional entry in our Hessian can be computed efficiently using Chebyshev approximations, making second order methods more efficient overall. We provide a lesion study comparison in Section 6.3.

### 4.3 Practical Implementation

In this section, we outline implementation concerns that are important for querying the moments sketch in practice. We include a number of optimizations to improve the stability and performance of Newton’s method, and also discuss the stability of the moments sketch under floating point precision. Due to space constraints, some equations are omitted and provided in Appendix A and B in [30].

#### 4.3.1 Optimizing Newton’s Method

The primary source of difficulties in implementing Newton’s method is the Hessian  $\nabla^2 L$  of our objective  $L$ . In our case:

$$\nabla^2 L(\theta)_{ij} = \int_{x_{\min}}^{x_{\max}} m_i(x)m_j(x)f(x;\theta) dx, \quad (6)$$

where the functions  $m_i(x)$  range over the set of functions

$$\{x^i : i \in \{1, \dots, k_1\}\} \cup \{\log^i(x) : i \in \{1, \dots, k_2\}\}.$$

There are two main challenges in performing a Newton step using this Hessian. First,  $\nabla^2 L$  can be nearly singular and cause numerical instabilities in Newton’s method that prevent or slow down convergence. Second, since the integral in Eq. (6) has no closed form, the cost of performing  $O(k^2)$  numerical integrations to compute  $\nabla^2 L$  in each iteration can be expensive.

**Conditioning.** To quantify the degree of numerical instability, we use the *condition number* of the Hessian  $\nabla^2 L$ . The condition number  $\kappa(A)$  of a matrix  $A$  describes how close a matrix is to being singular: matrices with high condition number are close to being singular, and  $\log_{10} \kappa$  provides an estimate of how many digits of precision are lost when inverting  $A$ . In particular, the use of the powers  $m_i(x) \in \{x^i : i \in \{1, \dots, k_1\}\}$  can result in ill-conditioned Hessians [31]. For example, when solving for a maximum entropy distribution with  $k_1 = 8, k_2 = 0, x_{\min} = 20$ , and  $x_{\max} = 100$ , we encountered  $\kappa(\nabla^2 L) \approx 3 \cdot 10^{31}$  at  $\theta = 0$ , making even the very first Newton step unstable.

We mitigate this issue by using a change of basis from the functions  $m_i(x) = x^j$  and  $m_i(x) = \log^j(x)$  to the basis of Chebyshev polynomials  $T_i(x)$ . Chebyshev polynomials are bounded polynomials supported on  $[-1, 1]$  and are often used for polynomial approximation [10, 64]. Using them we define the new basis  $\tilde{m}_i$  as follows:

$$\tilde{m}_i(x) = \begin{cases} T_i(s_1(x)), & i \in \{1, \dots, k_1\} \\ T_{i-k_1}(s_2(\log(x))), & i \in \{k_1 + 1, \dots, k_1 + k_2\} \end{cases}$$

where  $s_1, s_2$  are linear scaling functions that map to  $[-1, 1]$ . The new basis functions  $\tilde{m}_i(x)$  can be expressed in terms of  $x^j$  and  $\log^j(x)$  using standard formulae for Chebyshev polynomials and the binomial expansion [56]. Using this new basis for  $m_i$  Equation (6), we found that the condition number for the above example is reduced to  $\kappa \approx 11.3$ , making precision loss during each Newton step less of a concern.

On certain datasets, if ill-conditioned matrices are still an issue at query time we further limit ourselves to using the first  $k_1 \leq k$  moments and  $k_2 \leq k$  log moments by selecting  $k_1, k_2$  such that the condition number of the Hessian is less than a threshold  $\kappa_{\max}$ . Our heuristics select  $k_1, k_2$  by greedily incrementing  $k_1$  and  $k_2$  and favoring moments which are closer to the moments expected from a uniform distribution.

**Efficient Integration.** Naively computing the Hessian in Equation (6) requires evaluating  $O(k^2)$  numerical integrals per iteration, which can lead to prohibitively slow estimation time. We reduce this computational overhead by using polynomial approximations of the functions appearing in the integrands. If the integrands  $\tilde{m}_i(x)\tilde{m}_j(x)f(x;\theta)$  were expressible as polynomials in  $x$ , then each integral can be evaluated in closed form. The factors in the integrand that do not appear as polynomials in  $x$  are  $\tilde{m}_i(x), i \in \{k_1 + 1, \dots, k_1 + k_2\}$ , which are polynomials in  $\log(x)$ , and the pdf  $f(x;\theta)$ . Therefore, we compute Chebyshev polynomial approximations of these factors and replace each instance in the integrands with its corresponding approximation. For more details see Appendix A.2 in [30].

Approximating each of the factors with a degree  $n_c$  polynomial takes  $O(n_c \cdot \log n_c)$  using a fast cosine transform [64], so computing the Hessian can be done in  $O(k_2 n_c \log n_c + n_c k_1 k_2)$ . This is not an asymptotic improvement over naive numeric integration, but the number of complex function evaluations (i.e.  $\cos(x), e^x$ ) is reduced substantially. As we show in our empirical evaluation (Section 6.3), polynomial approximations reduce solve times  $20\times$  compared to numerically integrating each entry of the Hessian independently. We find in our experiments that the major bottleneck during maximum entropy optimization is the cosine transform used to construct the polynomial approximations.

#### 4.3.2 Floating point stability

Numeric floating point stability limits the range of useful  $k$  in a moments sketch. Both our estimation routine and error bounds (Section 4.4) use moments corresponding to data shifted onto the range  $[-1, 1]$ . On scaled data with range  $[c - 1, c + 1]$ , this leads to numeric error  $\epsilon_k$  in the  $k$ -th shifted moment, bounded by  $\epsilon_k \leq 2^k (|c| + 1)^k \epsilon_s$  where  $\epsilon_s$  is the relative error in the raw moments sketch power sums. This shift is the primary source of precision loss. We relate the loss to the error bounds in Section 4.4 to show that when using double precision floating point moments up to

$$k \leq \frac{13.06}{0.78 + \log_{10}(|c| + 1)} \quad (7)$$

provide numerically useful values. Data centered at 0 ( $c = 0$ ) have stable higher moments up to  $k = 16$ , and in practice we encounter issues when  $k \geq 16$ . We provide derivations and evaluations of this formula in Appendix B and C in [30]

### 4.4 Quantile Error Bounds

Recall that we estimate quantiles by constructing a maximum entropy distribution subject to the constraints recorded in a moments sketch. Since the true empirical distribution is in general not equal to the estimated maximum entropy distribution, to what extent can the quantiles estimated from the sketch deviate from the true quantiles? In this section, we discuss worst-case bounds on the discrepancy between *any* two distributions which share the same moments, and relate these to bounds on the quantile estimate errors. In practice, error on non-adversarial datasets is lower than these bounds suggest.

We consider distributions supported on  $[-1, 1]$ : we can scale and shift any distribution with bounded support to match. By Proposition 1 in Kong and Valiant [47], any two distributions supported on  $[-1, 1]$  with densities  $f$  and  $g$  and standard moments  $\mu_f, \mu_g$ , the Wasserstein distance

(or Earth Mover’s distance)  $W_1(f, g)$  between  $f$  and  $g$  is bounded by:

$$W_1(f, g) \leq O\left(\frac{1}{k} + 3^k \|\mu_f - \mu_g\|_2\right).$$

For univariate distributions  $f$  and  $g$ , the Wasserstein distance between the distributions is equal to the L1 distance between their respective cumulative distribution functions  $F$  and  $G$  (see Theorem 6.0.2 in [6]). Thus:

$$W_1(f, g) = \int_{-1}^{+1} |F(x) - G(x)| dx.$$

If  $f$  is the true dataset distribution, we estimate  $\hat{q}_\phi$  by calculating the  $\phi$ -quantile of the maximum entropy distribution  $\hat{f}$ . The quantile error  $\varepsilon(\hat{q}_\phi)$  is then equal to the gap between the CDFs:  $\varepsilon(q_\phi) = |F(\hat{q}_\phi) - \hat{F}(\hat{q}_\phi)|$ . Therefore, the average quantile error over the support  $[-1, 1]$  is bounded as follows:

$$\int_{-1}^{+1} \varepsilon(x) dx \leq O\left(\frac{1}{k} + 3^k \|\mu_f - \mu_{\hat{f}}\|_2\right). \quad (8)$$

Since we can run Newton’s method until the moments  $\mu_f$  and  $\mu_{\hat{f}}$  match to any desired precision, the  $3^k \|\mu_f - \mu_{\hat{f}}\|_2$  term is negligible.

Equation (8) does not directly apply to the  $\epsilon_{\text{avg}}$  used in Section 6, which is averaged over  $\phi$  for uniformly spaced  $\phi$ -quantiles rather than over the support of the distribution. Since  $\phi = \hat{F}(\hat{q}_\phi)$ , we can relate  $\epsilon_{\text{avg}}$  to Eq. (8) using our maximum entropy distribution  $\hat{f}$ :

$$\epsilon_{\text{avg}} = \int_0^1 \varepsilon(\hat{q}_\phi) d\phi = \int_{-1}^{+1} \varepsilon(x) \hat{f}(x) dx \leq O\left(\frac{\hat{f}_{\text{max}}}{k}\right)$$

where  $\hat{f}_{\text{max}}$  is the maximum density of our estimate. Thus, we expect the average quantile error  $\epsilon_{\text{avg}}$  to have a decreasing upper bound as  $k$  increases, with higher potential error when  $\hat{f}$  has regions of high density relative to its support. Though these bounds are too conservative to be useful in practice, they provide useful intuition on how worst case error can vary with  $k$  and  $\hat{f}$  (Figure 23).

## 5. THRESHOLD QUERIES

We described in Section 3.3 two types of queries: single quantile queries and threshold queries over multiple groups. The optimizations in Section 4.3 can bring quantile estimation overhead down to  $\leq 1\text{ms}$ , which is sufficient for interactive latencies on single quantile queries. In this section we show how we can further reduce quantile estimation overheads on threshold queries. Instead of computing the quantile on each sub-group directly, we compute a sequence of progressively more precise bounds in a *cascade* [75], and only use more expensive estimators when necessary. We first describe a series of bounds relevant to the moments sketch in Section 5.1 and then show how they can be used in end-to-end queries in Section 5.2.

### 5.1 Moment-based inequalities

Given the statistics in a moments sketch, we apply a variety of classical inequalities to derive bounds on the quantiles. These provide worst-case error guarantees for quantile estimates, and enable faster query processing for threshold queries over multiple groups.

---

### Algorithm 2: Threshold Query Cascade

---

```

macro CheckBound( $r_{\text{lower}}, r_{\text{upper}}, r_t$ )
  if  $r_{\text{lower}} > r_t$  then
    | return true
  else if  $r_{\text{upper}} < r_t$  then
    | return false
  function Threshold(threshold  $t$ , quantile  $\phi$ )
    if  $t > x_{\text{max}}$  then
      | return false
    if  $t < x_{\text{min}}$  then
      | return true
     $r_{\text{lower}}, r_{\text{upper}} \leftarrow$  MarkovBound( $t$ )  $\triangleright$  Markov Bound
    CheckBound( $r_{\text{lower}}, r_{\text{upper}}, n\phi$ )
     $r_{\text{lower}}, r_{\text{upper}} \leftarrow$  RTTBound( $t$ )  $\triangleright$  RTT Bound
    CheckBound( $r_{\text{lower}}, r_{\text{upper}}, n\phi$ )
     $q_\phi \leftarrow$  MaxEntQuantile( $\phi$ )  $\triangleright$  Maximum Entropy
    return  $q_\phi > t$ 

```

---

One simple inequality we make use of is Markov’s inequality. Given a non-negative dataset  $D$  with moments  $\mu_i$  Markov’s inequality tells us that for any value  $t$ ,  $\text{rank}(t) \geq n(1 - \frac{\mu_k}{t^k})$  where the rank is the number of elements in  $D$  less than  $t$ . We can apply Markov’s inequality to moments of transformations of  $D$  including  $T^+(D) = \{x - x_{\text{min}} : x \in D\}$ ,  $T^-(D) = \{x_{\text{max}} - x : x \in D\}$ , and  $T^l(D) = \{\log(x) : x \in D\}$  to bound  $\text{rank}(t)$  and thus also the error  $\epsilon$  for quantile estimates  $t = \hat{q}_\phi$ . We refer to this procedure as the MarkovBound procedure.

The authors in [66] provide a procedure (Section 3, Figure 1 in [66]) for computing tighter but more computationally expensive bounds on the CDF  $F(t)$  of a distribution given its moments. We refer to this procedure as the RTTBound procedure, and as with the MarkovBound procedure, use it to bound the error of a quantile estimate  $\hat{q}_\phi$ . The RTTBound procedure does not make use of the standard moments and log moments simultaneously, so we run RTTBound once on the standard moments and once on log moments and take the tighter of the bounds.

### 5.2 Cascades for Threshold queries

Given a moments sketch, Algorithm 2 shows how we calculate **Threshold**( $t, \phi$ ): whether the dataset has quantile estimate  $\hat{q}_\phi$  above a cutoff  $t$ . We use this routine to check whether a subgroup should be included in a query filter without computing  $\hat{q}_\phi$  directly. The threshold check routine first performs a simple filter on whether the threshold  $t$  falls in the range  $[x_{\text{min}}, x_{\text{max}}]$ . Then, we can use the Markov inequalities **MarkovBound** to calculate lower and upper bounds on the rank of the threshold  $\text{rank}(t)$  in the sub-population. Similarly the **RTTBound** routine uses more sophisticated inequalities in [66] to obtain tighter bounds on the rank. These bounds are used to determine if we can resolve the threshold predicate immediately. If not, we solve for the maximum entropy distribution as described in Section 4.2 (**MaxEntQuantile**) and calculate  $\hat{q}_\phi$ .

The Markov and RTTBound bounds are cheaper to compute than our maximum entropy estimate, allowing us to short-circuit many of the threshold predicates without explicitly estimating quantiles. The bounds apply to any distribution that matches the moments in a moments sketch, so this routine has no false negatives and is consistent with calculating the maximum entropy quantile estimate up front.

## 6. EVALUATION

In this section we evaluate the efficiency and accuracy of the moments sketch in a series of microbenchmarks, and then show how the moments sketch provides end-to-end performance improvements in the Druid and Macrobase data analytics engines [8, 82].

This evaluation demonstrates that:

1. The moments sketch supports 15 to 50× faster query times than comparably accurate summaries on quantile aggregations.
2. The moments sketch provides  $\epsilon_{\text{avg}} \leq 0.01$  estimates across a range of real-world datasets using less than 200 bytes of storage.
3. Maximum entropy estimation is more accurate than alternative moment-based quantile estimates, and our solver improves estimation time by 200× over naive solutions.
4. Integrating the moments sketch as a user-defined sketch provides 7× faster quantile queries than the default quantile summary in Druid workloads.
5. Cascades can provide 25× higher query throughput compared to direct moments sketch usage in Macrobase threshold queries.

Throughout the evaluations, the moments sketch is able to accelerate a variety of aggregation-heavy workloads with minimal space overhead.

### 6.1 Experimental Setup

We implement the moments sketch and its quantile estimation routines in Java<sup>1</sup>. This allows for direct comparisons with the open source quantile summaries [1, 67] and integration with the Java-based Druid [82] and MacroBase [8] systems. In our experimental results, we use the abbreviation M-Sketch to refer to the moments sketch.

We compare against a number of alternative quantile summaries: a mergeable equi-width histogram (**EW-Hist**) using power-of-two ranges [65], the ‘GKArray’ (**GK**) variant of the Greenwald Khanna [34, 52] sketch, the AVL-tree T-Digest (**T-Digest**) [28] sketch, the streaming histogram (**S-Hist**) in [12] as implemented in Druid, the ‘Random’ (**RandomW**) sketch from [52, 77], reservoir sampling (**Sampling**) [76], and the low discrepancy mergeable sketch (**Merge12**) from [3], both implemented in the Yahoo! datasketches library [1]. The **GK** sketch is not usually considered mergeable since its size can grow upon merging [3], this is especially dramatic in the production benchmarks in Appendix D.4 in [30]. We do not compare against fixed-universe quantile summaries such as the Q-Digest [71] or Count-Min sketch [23] since they would discretize continuous values.

Each quantile summary has a *size parameter* controlling its memory usage, which we will vary in our experiments. Our implementations and benchmarks use double precision floating point values. During moments sketch quantile estimation we run Newton’s method until the moments match to within  $\delta = 10^{-9}$ , and select  $k_1, k_2$  using a maximum condition number  $\kappa_{\text{max}} = 10^4$ . We construct the moments sketch to store both standard and log moments up to order  $k$ , but choose at query time which moments to make use of as described in Section 4.3.2.

<sup>1</sup><https://github.com/stanford-futuredata/msketch>

Table 1: Dataset Characteristics

	milan	hepmass	occupancy	retail	power	expon
size	81M	10.5M	20k	530k	2M	100M
min	$2.3e-6$	-1.961	412.8	1	0.076	$1.2e-7$
max	7936	4.378	2077	80995	11.12	16.30
mean	36.77	0.0163	690.6	10.66	1.092	1.000
stddev	103.5	1.004	311.2	156.8	1.057	0.999
skew	8.585	0.2946	1.654	460.1	1.786	1.994

We quantify the accuracy of a quantile estimate using the quantile error  $\epsilon$  as defined in Section 3.1. Then, as in [52, 77] we can compare the accuracies of summaries on a given dataset by computing their *average error*  $\epsilon_{\text{avg}}$  over a set of uniformly spaced  $\phi$ -quantiles. In the evaluation that follows, we test on 21 equally spaced  $\phi$  between 0.01 and 0.99.

We evaluate each summary via single-threaded experiments on a machine with an Intel Xeon E5-4657L 2.40GHz processor and 1TB of RAM, omitting the time to load data from disk.

#### 6.1.1 Datasets

We make use of six real-valued datasets in our experiments, whose characteristics are summarized in Table 1. The *milan* dataset consists of Internet usage measurements from Nov. 2013 in the Telecom Italia Call Data Records [40]. The *hepmass* dataset consists of the first feature in the UCI [49] HEPMASS dataset. The *occupancy* dataset consists of CO2 measurements from the UCI Occupancy Detection dataset. The *retail* dataset consists of integer purchase quantities from the UCI Online Retail dataset. The *power* dataset consists of Global Active Power measurements from the UCI Individual Household Electric Power Consumption dataset. The *exponential* dataset consists of synthetic values from an exponential distribution with  $\lambda = 1$ .

### 6.2 Performance Benchmarks

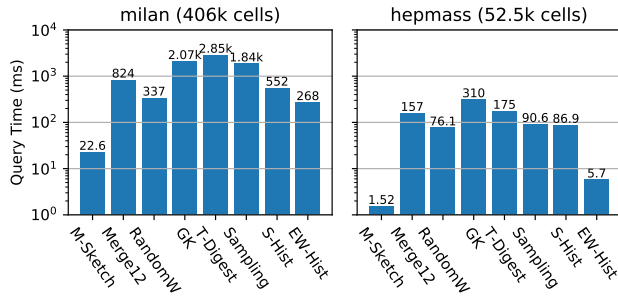
We begin with a series of microbenchmarks evaluating the moments sketch query times and accuracy.

#### 6.2.1 Query Time

Our primary metric for evaluating the moments sketch is total query time. We evaluate quantile aggregation query times by pre-aggregating our datasets into cells of 200 values and maintaining quantile summaries for each cell. Then we measure the time it takes to performing a sequence of merges and estimate a quantile. In this performance microbenchmark, the cells are grouped based on their sequence in the dataset, while the evaluations in Section 7 group based on column values. We divide the datasets into a large number of cells to simulate production data cubes, while in Appendix D.3 and D.4 in [30] we vary the cell sizes. Since the moments sketch size and merge time are data-independent, the results generalize as we vary cell size.

Figure 3 shows the total query time to merge the summaries and then compute a quantile estimate when each summary is instantiated at the smallest size sufficient to achieve  $\epsilon_{\text{avg}} \leq .01$  accuracy. We provide the parameters we used and average observed space usage in Table 2. On the long-tailed milan dataset, the **S-Hist** and **EW-Hist** summaries are unable to achieve  $\epsilon_{\text{avg}} \leq .01$  accuracy with less than 100 thousand buckets, so we provide timings at 100 buckets for comparison. The moments sketch provides 15





**Figure 3:** Total query time using different summaries to estimate quantiles with  $\epsilon_{\text{avg}} \leq .01$ . The moments sketch enables significantly faster queries at this accuracy.

**Table 2:** Summary size parameters for  $\epsilon_{\text{avg}} \leq .01$ .

dataset	milan	size (b)	hepmass	size (b)
sketch	param		param	
M-Sketch	$k = 10$	200	$k = 3$	72
Merge12	$k = 32$	5920	$k = 32$	5150
RandomW	$\epsilon = \frac{1}{40}$	3200	$\epsilon = \frac{1}{40}$	3375
GK	$\epsilon = \frac{1}{60}$	720	$\epsilon = \frac{1}{40}$	496
T-Digest	$\delta = 5.0$	769	$\delta = 1.5$	93
Sampling	1000 samples	8010	1000	8010
S-Hist	100 bins	1220	100	1220
EW-Hist	100 bins	812	15	132

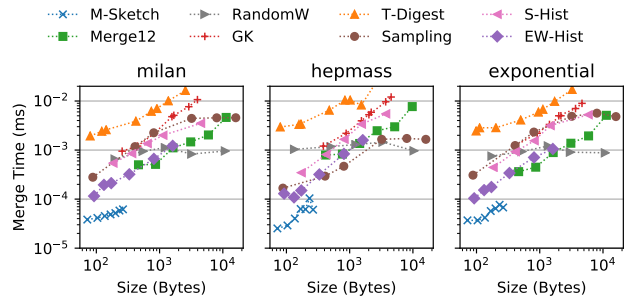
to  $50\times$  faster query times than **RandomW**, the next fastest accurate summary. As a baseline, sorting the milan dataset takes 7.0 seconds, selecting an exact quantile online takes 880ms, and streaming the data pointwise into a **RandomW** sketch with  $\epsilon = 1/40$  takes 220ms. These methods do not scale as well as using pre-aggregated moments sketches as dataset density grows but the number of cells remains fixed.

### 6.2.2 Merge and Estimation Time

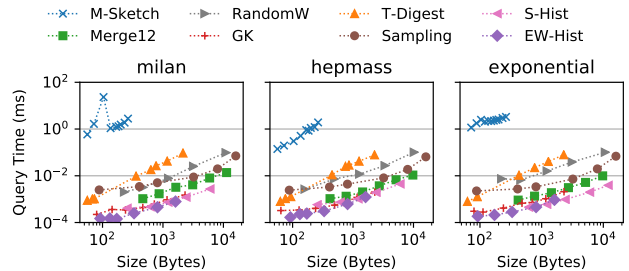
Recall that for a basic aggregate quantile query  $t_{\text{query}} = t_{\text{merge}} \cdot n_{\text{merge}} + t_{\text{est}}$ . Thus we also measure  $t_{\text{merge}}$  and  $t_{\text{est}}$  to quantify the regimes where the moments sketch performs well. In these experiments, we vary the summary size parameters, though many summaries have a minimum size, and the moments sketch runs into numeric stability issues past  $k \geq 15$  on some datasets (see Section 4.3.2).

In Figure 4 we evaluate the average time required to merge one of the cell summaries. Larger summaries are more expensive to merge, and the moments sketch has faster ( $< 50\text{ns}$ ) merge times throughout its size range. When comparing summaries using the parameters in Table 2, the moments sketch has up to  $50\times$  faster merge times than other summaries with the same accuracy.

One can also parallelize the merges by sharding the data and having separate nodes operate over each partition, generating partial summaries to be aggregated into a final result. Since each parallel worker can operate independently, in these settings the moments sketch maintains the same relative performance improvements over alternative summaries when we can amortize fixed overheads, and we include supplemental parallel experiments in Appendix F in [30]. The other major contributor to query time is estimation time. In Figure 5 we measure the time to estimate quantiles given an



**Figure 4:** Per-merge latencies. The moments sketch provides faster merge times than alternative summaries at the same size.



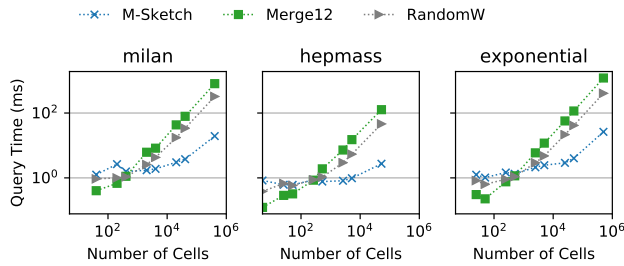
**Figure 5:** Quantile Estimation time. Estimation time on the moments sketch is slower than other sketches but under 3ms for  $k = 10$ .

existing summary. The moments sketch provides on average 2 ms estimation times, though estimation time can be higher when our estimator chooses higher  $k_1, k_2$  to achieve better accuracy. This is the cause for the spike at  $k = 4$  in the milan dataset and users can mitigate this by lowering the condition number threshold  $\kappa_{\text{max}}$ . Other summaries support microsecond estimation times. The moments sketch thus offers a tradeoff of better merge time for worse estimation time. If users require faster estimation times, the cascades in Section 5.2 and the alternative estimators in Section 6.3 can assist. We show how the merge time and estimation time tradeoff define regimes where each component dominates depending on the number of merges. In Figure 6 we measure how the query time changes as we vary the number of summaries (cells of size 200) we aggregate. We use the moments sketch with  $k = 10$  and compare against the mergeable **Merge12** and **RandomW** summaries with parameters from Table 2. When  $n_{\text{merge}} \geq 10^4$  merge time dominates and the moments sketch provides better performance than alternative summaries. However, the moments sketch estimation times dominate when  $n_{\text{merge}} \leq 100$ .

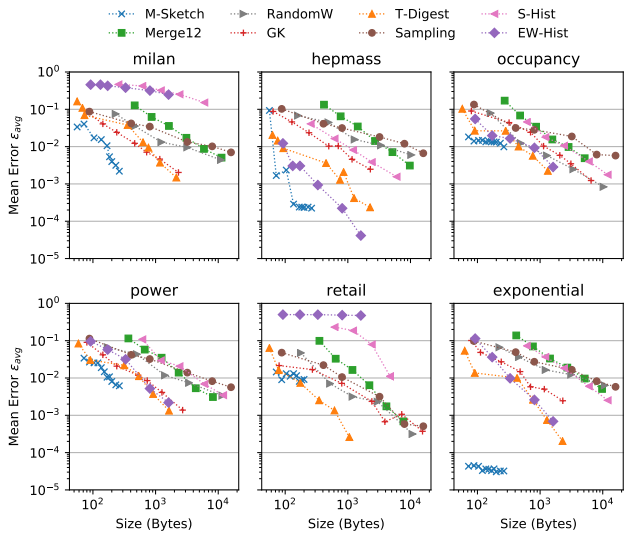
### 6.2.3 Accuracy

The moments sketch accuracy is dataset dependent, so in this section we compare the average quantile error on our evaluation datasets.

Figure 7 illustrates the average quantile error  $\epsilon_{\text{avg}}$  for summaries of different sizes constructed using pointwise accumulation on the complete dataset. The moments sketch achieves  $\epsilon \leq 10^{-4}$  accuracy on the synthetic exponential dataset, and  $\epsilon \leq 10^{-3}$  accuracy on the high entropy hepmass



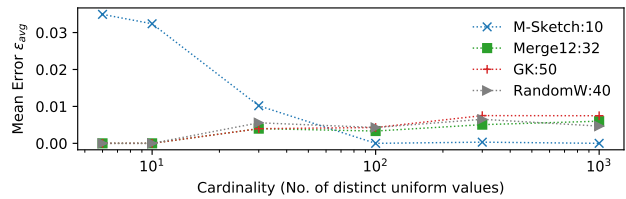
**Figure 6:** Comparing total query time using different mergeable summaries as we vary the number of merges. The moments sketch provides performance improvements when  $n_{\text{merge}} \geq 10^4$ .



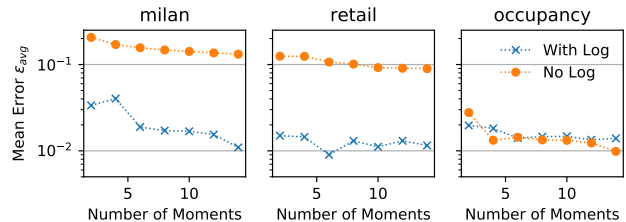
**Figure 7:** Average error for summaries of different sizes. The moments sketch delivers consistent  $\epsilon_{\text{avg}} \leq 0.015$  with fewer than 200 bytes.

dataset. On other datasets it is able to achieve  $\epsilon_{\text{avg}} \leq 0.01$  with fewer than 200 bytes of space. On the integer retail dataset we round estimates to the nearest integer. The EW-Hist summary, while efficient to merge, provides less accurate estimates than the moments sketch, especially in the long-tailed milan and retail datasets.

We provide further experiments in [30] showing how the moments sketch worst-case error bounds are comparable to other summaries (Appendix E), that the moments sketch is robust to changes in skew and the presence of outliers (Appendix D), and that the moments sketch generalizes to a production workload (Appendix D.4). However, on datasets with low-entropy, in particular datasets consisting of a small number of discrete point masses, the maximum entropy principle provides poor accuracy. In the worst case, the maximum entropy solver can fail to converge on datasets with too few distinct values. Figure 8 illustrates how the error of the maximum entropy estimate increases as we lower the cardinality of a dataset consisting of uniformly spaced points in the range  $[-1, 1]$ , eventually failing to converge on datasets with fewer than five distinct values. If users are expecting to run queries on primarily low-cardinality datasets, fixed-



**Figure 8:** Accuracy of maximum entropy estimates on distributions with varying cardinality. The moments sketch is less accurate on discretized datasets, and fails to converge for cardinalities  $n < 5$ .



**Figure 9:** Accuracy with and without log moments. Given the same total space budget, log moments improve accuracy on the long-tailed milan and retail datasets, and do not affect accuracy significantly on other datasets such as occupancy

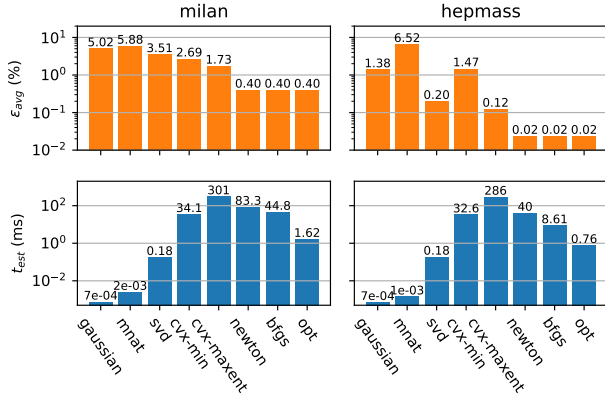
universe sketches or heavy-hitters sketches may be more appropriate.

### 6.3 Quantile Estimation Lesion Study

To evaluate each component of our quantile estimator design, we compare the accuracy and estimation time of a variety of alternative techniques on the milan and hepmass datasets. We evaluate the impact of using log moments, the maximum entropy distribution, and our optimizations to estimation.

To examine effectiveness of log moments, we compare our maximum entropy quantile estimator accuracy with and without log moments. For a fair comparison, we compare the estimates produced from  $k$  standard moments and no log moments with those produced from up to  $\frac{k}{2}$  of each. Figure 9 illustrates how on some long-tailed datasets, notably milan and retail, log moments reduce the error from  $\epsilon > .15$  to  $\epsilon < .015$ . On other datasets, log moments do not have a significant impact.

We compare our estimator (`opt`) with a number of other estimators that make use of the same moments. The gaussian estimator fits a Gaussian distribution to the mean and standard deviation. The `mnat` estimator uses the closed form discrete CDF estimator in [58]. The `svd` estimator discretizes the domain and uses singular value decomposition to solve for a distribution with matching moments. The `cvx-min` estimator also discretizes the domain and uses a convex solver to construct a distribution with minimal maximum density and matching moments. The `cvx-maxent` estimator discretizes the domain and uses a convex solver to maximize the entropy, as described in Chapter 7 in [15]. The `newton` estimator implements our estimator without the integration techniques in Sec. 4.3, and uses adaptive Romberg inte-



**Figure 10:** Lesion study comparing our optimized maximum entropy solver to other estimators. Our `opt` estimator provides at least  $5\times$  less error than estimators that do not use maximum entropy, and up to  $200\times$  faster estimation times than naive maximum entropy solvers.

gration instead [64]. The `bfgs` estimator implements maximum entropy optimization using the first-order L-BFGS [50] method as implemented in a Java port of `liblbfgs` [46].

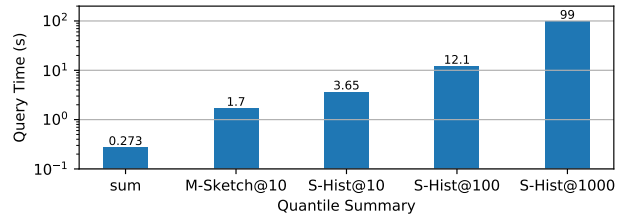
Figure 10 illustrates the average quantile error and estimation time for these estimators. We run these experiments with  $k = 10$  moments. For uniform comparisons with other estimators, on the milan dataset we only use the log moments, and on the hepmass dataset we only use the standard moments. We perform discretizations using 1000 uniformly spaced points and make use of the ECOS convex solver [27]. Solvers that use the maximum entropy principle provides at least  $5\times$  less error than estimators that do not. Furthermore, our optimizations are able to improve the estimation time by a factor of up to  $200\times$  over an implementation using generic solvers, and provide faster solve times than naive Newton’s method or BFGS optimizers. As described in Section 4.3, given the computations needed to calculate the gradient, one can compute the Hessian relatively cheaply, so our optimized Newton’s method is faster than BFGS.

## 7. APPLYING THE MOMENTS SKETCH

In this section, we evaluate how the moments sketch affects performance when integrated with other data systems. We examine how the moments sketch improves query performance in the Druid analytics engine, as part of a cascade in the Macrobase feature selection engine [8], and as part of exploratory sliding window queries.

### 7.1 Druid Integration

To illustrate the utility of the moments sketch in a modern analytics engine, we integrate the moments sketch with Druid [82]. We do this by implementing moments sketch as an user-defined aggregation extension, and compare the total query time on quantile queries using the moments sketch with the default `S-Hist` summary used in Druid and introduced in [12]. The authors in [12] observe on average 5% error for an `S-Hist` with 100 centroids, so we benchmark a moments sketch with  $k = 10$  against `S-Hists` with 10, 100, and 1000 centroids.



**Figure 11:** Druid end-to-end query benchmark. The moments sketch allows for faster query times than the comparable `S-Hist` summary with 100 bins. Runtime for a native sum operation is a lower bound on query time.

In our experiments, we deploy Druid on a single node – the same machine described in section 6.1 – with the same base configuration used in the default Druid quickstart. In particular, this configuration dedicates 2 threads to process aggregations. Then, we ingest 26 million entries from the milan dataset at a one hour granularity and construct a cube over the grid ID and country dimensions, resulting in 10 million cells.

Figure 11 compares the total time to query for a quantile on the complete dataset using the different summaries. The moments sketch provides  $7\times$  lower query times than a `S-Hist` with 100 bins. Furthermore, as discussed in Section 6.2.1, any `S-Hist` with fewer than 10 thousand buckets provides worse accuracy on milan data than the moments sketch. As a best-case baseline, we also show the time taken to compute a native sum query on the same data. The 1 ms cost of solving for quantile estimates from the moments sketch on this dataset is negligible here.

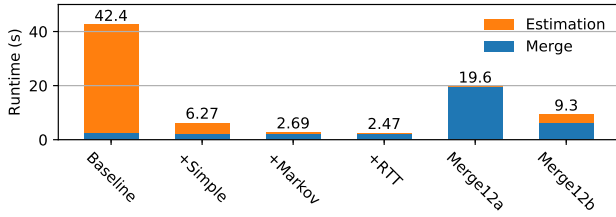
## 7.2 Threshold queries

In this section we evaluate how the cascades described in Section 5.2 improve performance on threshold predicates. First we show in Section 7.2.1 how the MacroBase analytics engine can use the moments sketch to search for anomalous dimension values. Then, we show in Section 7.2.2 how historical analytics queries can use the moments sketch to search and alert on sliding windows.

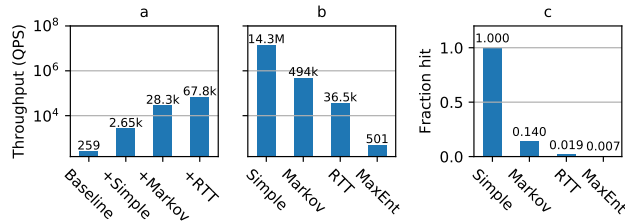
### 7.2.1 MacroBase Integration

The MacroBase engine searches for dimension values with unusually high outlier rates in a dataset [8]. For example, given an overall 2% outlier rate, MacroBase may report when a specific app version has an outlier rate of 20%. We integrate the moments sketch with a simplified deployment of MacroBase where all values greater than the global 99th percentile  $t_{99}$  are considered outliers. We then query MacroBase for all dimension values with outlier rate at least  $r = 30\times$  greater than the overall outlier rate. This is equivalent to finding subpopulations whose 70th percentile is greater than  $t_{99}$ .

Given a cube with pre-aggregated moments sketches for each dimension value combination and no materialized roll-ups, MacroBase merges the moments sketches to calculate the global  $t_{99}$ , and then runs Algorithm 2 on every dimension-value subpopulation, searching for subgroups with  $q_{.7} > t_{99}$ . We evaluate the performance of this query on 80 million rows of the milan internet usage data from November 2013,



**Figure 12:** Runtime of MacroBase queries: the final moments sketch cascade outperforms alternate sketches.



**Figure 13:** Cascades in MacroBase: (a) as we add stages, query throughput increases. (b) The cascade proceeds from faster to slower estimates. (c) Each stage processes a smaller fraction of queries.

pre-aggregated by grid ID, country, and at a four hour granularity. This resulted in 13 million cube cells, each with its own moments sketch.

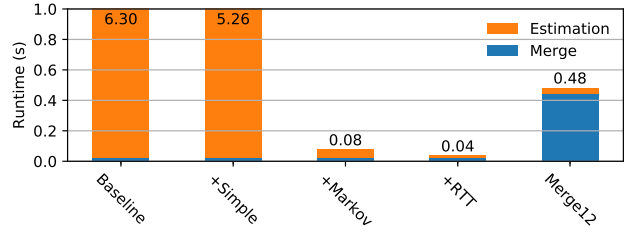
Running the MacroBase query produces 19 candidate dimension values. We compare the total time to process this query using direct quantile estimates, our cascades, and the alternative `Merge12` quantile sketch. In the first approach (`Merge12a`), we merge summaries during MacroBase execution as we do with a moments sketch. In the second approach (`Merge12b`), we calculate the number of values greater than the  $t_{99}$  for each dimension value combination and accumulate these counts directly, instead of the sketches. We present this as an optimistic baseline, and is not always a feasible substitute for merging summaries.

Figure 12 shows the query times for these different methods: the baseline method calculates quantile estimates directly, we show the effect of incrementally adding each stage of our cascade ending with `+RTTBound`. Each successive stage of the cascade improves query time substantially. With the complete cascade, estimation time is negligible compared to merge time. Furthermore, the moments sketch with cascades has  $7.9\times$  lower query times than using the `Merge12` sketch, and even  $3.7\times$  lower query times than the `Merge12b` baseline.

In Figure 13 we examine the impact the cascade has on estimation time directly. Each additional cascade stage improves threshold query throughput and is more expensive than the last. The complete cascade is over  $250\times$  faster than this baseline, and  $25\times$  faster than just using a simple range check.

### 7.2.2 Sliding Window Queries

Threshold predicates are broadly applicable in data exploration queries. In this section, we evaluate how the moments sketch performs on sliding window alerting queries. This is



**Figure 14:** Sliding window query: moments sketch with cascades runs  $13\times$  faster than `Merge12`.

useful when, for instance, users are searching for time windows of unusually high CPU usage spikes.

For this benchmark, we aggregated the 80 million rows of the milan dataset at a 10-minute granularity, which produced 4320 panes that spanned the month of November. We augmented the milan data with two spikes corresponding to hypothetical anomalies. Each spike spanned a two-hour time frame and contributed 10% more data to those time frames. Given a global 99th percentile of around 500 and a maximum value of 8000, we added spikes with values  $x = 2000$  and  $x = 1000$

We then queried for the 4-hour time windows whose 99th percentile was above a threshold  $t = 1500$ . When processing this query using a moments sketch, we can update sliding windows using turnstile semantics, subtracting the values from the oldest pane and merging in the new one, and use our cascade to filter windows with quantiles above the threshold.

Figure 14 shows the runtime of the sliding window query using both the moments sketch and `Merge12`. Faster moments sketch merge times and the use of turnstile semantics then allow for  $13\times$  faster queries than `Merge12`.

## 8. CONCLUSION

In this paper, we show how to improve the performance of quantile aggregation queries using statistical moments. Low merge overhead allows the moments sketch to outperform comparably accurate existing summaries when queries aggregate more than 10 thousand summaries. By making use of the method of moments and the maximum entropy principle, the moments sketch provides  $\epsilon_{\text{avg}} \leq 0.01$  accuracy on real-world datasets, while the use of numeric optimizations and cascades keep query times at interactive latencies.

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