Abstract

Given some of the recent advances in Distributed Hash Table (DHT) based Peer-To-Peer (P2P) systems we ask the following questions: Are there applications where unstructured queries are still necessary (i.e., the underlying queries do not efficiently map onto any structured framework), and are there unstructured P2P systems that can deliver the high bandwidth and computing performance necessary to support such applications. Toward this end, we consider an image search application which supports queries based on image similarity metrics, such as color histogram intersection, and discuss why in this setting, standard DHT approaches are not directly applicable. We then study the feasibility of implementing such an image search system on two different unstructured P2P systems: power-law topology with percolation search, and an optimized super-node topology using structured broadcasts. We examine the average and maximum values for node bandwidth, storage and processing requirements in the percolation and super-node models, and show that current high-end computers and high-speed links have sufficient resources to enable deployments of large-scale complex image search systems.

1 Introduction

The first widely known pure P2P system that tried to bring Napster-like functionality was the unstructured P2P system Gnutella. The overlay network created by Gnutella’s peers forms a random graph, where search was mostly done via complete flooding of the network. Its imperfections have spawned extensive research. Much of this research is directed towards DHTs (e.g. [5, 12]), and distributed indexing structures derived from DHTs (e.g. [13]), super peer architectures (e.g. [15]) and approaches improving the link structure of P2P networks (e.g. [7]).

DHTs excel at key-value lookup because that is the basic primitive of the hash table data structure. However, hash tables are not the most efficient data structure for all algorithms. In this work we are interested in storing images, which may be thought of as vectors of large dimension, and searching to find images which are close to a query image by some given metric. This work compares content based image retrieval (CBIR) in structured against unstructured P2P systems. Unstructured systems can answer completely general queries, since there is no structure imposed on the data. We ask if sacrificing the flexibility of unstructured systems for structured P2P systems results in a reduction of query bandwidth for the case of CBIR.

Within this article, we treat content-based image retrieval as an application scenario. We think that this scenario is interesting for P2P in several respects. (1) We believe that there is a need for such applications. The recent success of image blogs and image sharing servers like flickr.com has shown that people have the wish to share and to publish their images. (2) Current widespread methods of indexing such images are unsatisfactory. In flickr.com, the images are searchable by annotation. Unfortunately the annotation quality is low (as will be described more in-depth below). Orga-
nizing images by time [2] is useful when indexing one’s own collection where images can usually be grouped into images pertaining to events that are interesting to the user, but it becomes barely useful when considering collections that grow each second by at least one image. (3) Alternative methods are expensive. Sophisticated content-based image indexing methods that extract visual features from the items to be queried need a large amount of processing power, and processing queries is expensive\(^1\) compared to processing text queries.

While CBIR is deemed unsatisfactory as a complete indexing solution for images, the conviction underlying this paper is that CBIR methods are going to be useful for improving \texttt{flickr.com} like systems. Bad image annotation is not going to go away. So, we seek systems that help the user in the case that good annotation is not present. We are also convinced that people will be interested in \textit{combined rankings}, which combine visual aspects, surrounding text, etc. into one common measure. Unless one wants to restrict oneself to systems that first evaluate similarity with respect to text and simple meta-data before refining the search using CBIR and other complex methods, one will need to come up with good methods that are able to process CBIR queries.

This paper presents two findings. First, we find that unstructured systems are efficient enough in terms of communication and computational costs for the CBIR application we consider to scale to millions of users. Second, in the case of image similarity search, current structured systems offer no advantage over unstructured systems. This paper is organized as follows. Section 2 discusses prior work on the nearest-neighbor search problem in high dimensional spaces, a general version of the problem we consider in this work. Section 3 describes the particular case we are focusing on: image similarity search at the scale of approximately one million users. Section 4.1 studies the cost of implementing an image search system using a supernode architecture and Section 4.2 studies the cost for a percolation search based architecture. Finally, in Section 5 we compare the unstructured image search systems to a structured image search system and find that structured search offers no benefit for this case.

2 Nearest neighbor search in high dimension

The nearest-neighbor search problem is the following: given a set of points in a metric space \(P\), for a given query point \(Q\), find an element \(x\) of \(P\) such that \(d(x, Q) \leq d(y, Q)\) for all \(y \in P\).

The classic way of performing CBIR is to extract real-valued feature vectors from images and then map the search problem to the problem of finding the \(k\) nearest neighbors (\(k\)-NN) to the query vector. For \(d <\approx 10\) there exist centralized data structures that find the \(k\)-NN in \(O(\log N)\) time for collections of size \(N\). However, literature on non-distributed indexing has observed [14] that \textit{exact} search in high-dimensional data is very hard, due to the so-called \textit{curse of dimensionality}. Due to the curse of dimensionality tree-based indexing structures break down in the sense that in realistic scenarios \(O(N)\) nodes need to be visited before finding the \textit{exact} \(k\)-NN. For non-distributed —disk based— indexing structures one interesting and well-known solution [14] consists in rendering full scan queries more efficient using a full table scan approach. This algorithm exhibits \(O(N)\) complexity just as a tree-based indexing structure in high dimensions, however, the absolute query duration is reduced with respect to the tree-based solution.

Though not the subject of this work, one may also consider the approximate version of the nearest-neighbor search problem. The approximate nearest-neighbor search problem is the following: given a set of points in a metric space \(P\), for a given query point \(Q\), and a slackness parameter \(\epsilon\) find an element \(x\) of \(P\) such that \(d(x, Q) \leq (1 + \epsilon)d(y, Q)\) for all \(y \in P\). An efficient algorithm for the approximate nearest-neighbor search problem is known for several common metric spaces [4]. It is an interesting open problem to see if it can be efficiently adapted to a distributed system.

Other proposals include using geometric dimensionality reduction techniques. Kleis and Zhou provide a review of relevant results in the context of P2P networks in [3].

2.1 P2P approaches to nearest-neighbor search

Clearly, finding the set \(
\{x|d(x, Q) \leq \delta\}
\) for a given \(Q, \delta\) is an embarrassingly parallel problem. If one spreads the database over a P2P network, one gets the full benefit of parallelization. In the absence of an efficient exact algorithm for the nearest-neighbor problem in high dimension, this may be the best one can do. Indeed we consider this approach in Sections 4.1 and 4.2.

In addition to the above, one can also attempt to use some structured P2P network to reduce the number of nodes that must be contacted to execute a query. We describe one such approach, PRISM below. The literature on P2P indexing using structured networks

\(^1\)That means, they typically require many disk accesses, much processing power and as a consequence much time to be processed.
is very broad. As one starting point for reading we suggest [13].

2.1.1 PRISM

PRISM indexes each vector \( \mathbf{x} \) by placing \( \mathbf{x} \) on a small number of nodes in a Chord DHT. The placement of the vector is calculated using distances to a fixed set of reference vectors. When processing a query, the node issuing the query \( q \) calculates the set of nodes where \( q \) would be placed and searches for similar nodes there, sending them \( q \) as the query. The main innovation of PRISM is the algorithm for finding the nodes on which to place the data vectors.

In order to index a vector \( \mathbf{x} \), the distance of \( \mathbf{x} \) to a number \( n_r \) of reference vectors \( r_i \) is calculated, yielding \( \delta := \{ \delta(\mathbf{x},r_1), \ldots, \delta(\mathbf{x},r_n) \} \). Then the \( r_i \) are ranked by their similarity. The result of this ranking is a list of indices \( \tau = (\tau_1, \ldots, \tau_{n_r}) \) such that \( r_{\tau_1} \) is the reference vector closest to \( \mathbf{x}, r_{\tau_2} \) the second closest and so on.

Then, pairs of indices are formed. The pair formation is a fitting parameter, the original PRISM paper suggests \( \{\tau_1, \tau_2\}, \{\tau_2, \tau_3\}, \{\tau_1, \tau_3\}, \{\tau_1, \tau_4\}, \{\tau_2, \tau_5\}, \{\tau_2, \tau_4\}, \{\tau_3, \tau_4\}, \{\tau_1, \tau_5\}, \{\tau_4, \tau_5\}, \{\tau_3, \tau_5\} \) for their dataset. From each of the pairs a Chord key is calculated, and this key is used for inserting the vector \( \mathbf{x} \) into the Chord ring.

As was hinted above, query processing works by finding out which peers would receive the query vector if it was a new data item and forwarding the query vector to these peers. This involves, again, the calculation of index pairs, which we will call query pairs in the following. In order to reduce query processing cost, the query processor can choose to contact only nodes pertaining to only a subset of the query pairs. Doing this also reduces recall, so there is a tradeoff.

3 An image search system

Within this section, we describe Flickr, a popular web-based photo sharing application. This application is currently immensely popular. At the same time, one could easily imagine extending its functionality towards content-based search. By examining Flickr we estimate the load for a P2P photo sharing system which we call Plickr.

3.1 About Flickr

flickr.com gives members the opportunity to share photos among the public, friends and family. Flickr members are allowed to comment on photos they can see and to annotate them in a collaborative fashion. Recently, flickr.com has experienced explosive growth of popularity. As of the time of writing, flickr.com contains about 40 million images, most of them publicly accessible. We estimate that about 2 million users share images via flickr.com.

One of the reasons for our interest into flickr.com is that it has a SOAP-like API that allows easy access. This simplifies enormously building third party tools, as well as crawlers. As the user structure is quite similar to what many people would like to have in P2P file sharing systems (people peacefully sharing data they actually own) we simply extrapolate from user behavior on flickr.com to the behavior they would have in a P2P network.

Using flickr.com, we do not obtain data about the online times of the users. However, information about who shares how much is already useful.

One of the most interesting features of Flickr’s is that members can annotate other people’s images. This leads to the surprising fact that most of Flickr’s images are annotated. However, as it is made simple to add annotation to images by default (on a user-by-user basis), the quality of the annotation is varying. For example, many Flickr members have a large fraction of their photos annotated with the tag phone. This tag describes how the image came to Flickr, by a camera built into a cellular phone. However, it is only rarely an accurate description of the images’ content, as the examples in Fig. 1 show. While the tag phone is clearly the most extreme case of annotation that carries little valuable information, it shows that just citing the number of images that are annotated does not permit assessing the usefulness of this annotation.

3.2 Plickr: content-based Flickr over P2P as a scenario for P2P-CBIR evaluation

The above ad-hoc assessment of annotation quality motivates the view that it would be interesting to combine the search by annotation tag (as offered by Flickr) by search based on image similarity as provided
by Content Based Image Retrieval systems (CBIRS). While CBIRS are unable to do true object recognition, they capture visual similarity by translating each image into a data representation that captures mainly image statistics and in some cases the spatial relation of “interesting” regions. While using such features as the sole image retrieval method is currently deemed unsatisfactory, CBIRS are still the method of choice when no annotation is present.

Content Based Image Retrieval is computationally costly. Features need to be extracted, and similarity search is much harder than similarity search on text because the number of features taken into account for obtaining a retrieval result is usually much higher than the number of keywords in a keyword query. For more information about CBIR we point to an often-cited overview paper [10].

P2P becomes particularly interesting through the fact that P2P-CBIR potentially will make use of both the huge storage capacity and the huge computing capacity distributed in the network.

Let us consider a P2P network, in which each peer owner shares his or her own photos with other users of the same P2P network. The network would offer Flickr’s functionality plus CBIR query by example functionality. However, in contrast to Flickr, its inner workings would be based entirely on P2P principles. We will call this hypothetical network the plickr network within this paper.

3.3 Deriving a load scenario for plickr

We assume the images contained in one Flickr user account to be a good model for the images shared by one plickr peer. Let us assume 1,000,000 users and thus 1,000,000 peers in our plickr network. Our Flickr crawls (≈2,000,000 images) indicate that the average user who shares at least one image publicly shares on average 20 images.

The same as Flickr, plickr members query the data collection for interesting images from time to time. Furthermore, we assume each user performs 10 queries on average per day. We feel that is reasonable as querying image is an exploratory process, so each querier is likely to perform a query process consisting of multiple queries. So, even if such a query process is performed less than once per day by each user, we are likely to reach the said average load.

In this paper we will concentrate on a very simple way of performing CBIR: retrieval by color histograms. They are known to provide a good retrieval performance (i.e. result quality) for comparatively little computing power and are the pet feature extraction method for indexing structure evaluations.

Color histograms are obtained by cutting the color space in regions. The color histogram then is a vector that contains one value corresponding to each color space region. Each value of the histogram expresses for the corresponding color region the probability that a pixel drawn from the image falls into the color region. This probability is estimated by simply counting pixels falling in each color region. The usefulness of color histograms for CBIR depends on the color space chosen and the way it is split into regions. For our load assumption, we assume John R. Smith’s 166-D HSV histograms described in [11]. A wasteful but simple representation would be of type float[166].

The classic way of evaluating the similarity of two histograms is the histogram intersection, however, the testing ground for most CBIR indexing algorithm is the Euclidean distance which is why we focus that distance measure.

4 Search with unstructured P2P

Unstructured P2P systems have one major advantage over structured systems: once designed, implemented and deployed an unstructured system can generally be used for any kind of query just by plugging in new query processing. There is no need to define new routing algorithms, network topologies, or caching strategies every time a new type of data or query is introduced to the network. On the other hand, structured P2P systems may reduce search complexity only if the search algorithm can be efficiently mapped onto the topology of the structured network.

In this section, we compute the costs in bandwidth, computational resources, and storage to use unstructured P2P for two models: a super-node system, and a percolation search system. After computing the costs of the system, in order to estimate feasibility, we make some assumptions about the usage of the image search system described in Section 3. We assumed that each content item is a float[166] array, which uses $166 \times 4 = 664$ bytes of space. As we mentioned in Section 3.3, every Flickr user inserts on average $C = 20$ items into the network. Calculating the distance takes $f = 332$ floating point operations. Finally, we will assume that there are $N = 2^{19} \approx 500,000$ users, and that each user will make 10 queries per day or $R = \frac{10}{2^{19} \times 60} = 1.2 \times 10^{-4}$ queries per second. We assume each query and content requires $z = 800$ bytes (enough to hold the float vector and some routing information or image meta-data).
Table 1. Nomenclature used throughout the paper

| Symbol | Meaning |
|--------|---------|
| \( N \) | Total number of peers |
| \( (k) \) | Expected degree of nodes within network |
| \( p_m \) | Relative frequency of nodes with degree \( m \) |
| \( s \) | Fraction of super peers |
| \( R \) | Query rate issued per peer (1.2 × 10^{-4} s^{-1}) |
| \( C \) | Number of content items contributed per peer (332 FLOPs) |
| \( f \) | Number of operations to compare two vectors |
| \( z \) | Size of each vector (800 B) |
| \( B_{\text{max,ave}} \) | Max/avg bandwidth required per peer |
| \( P_{\text{max,ave}} \) | Max/avg processing required per peer |
| \( D_{\text{max,ave}} \) | Max/avg disk space required per peer |

4.1 Super-node P2P networks

In the super-node model there are two types of nodes: leaf nodes and super-nodes. Each leaf node connects to a super-node and caches all its content on that super-node (the super-node does not need to cache its own content). The leaf nodes require very little resources, but super-nodes incur the maximum penalty. The only parameter in the system is \( s \), the fraction of nodes which are super-nodes.

4.1.1 Resource requirements

To compute average bandwidth, we count total copies of the queries and divide by the total number of nodes. We should note that this metric is not very meaningful since no nodes see the average. Leaf nodes see almost no traffic, while super-nodes see the maximum traffic.

Since each query is copied to \( sN \) super-nodes plus the leaf node that initiated the query, the average bandwidth is clearly, \( B_{\text{ave}} = \frac{R(N(sN+1))z}{N} = RNz(s+z^{-1}) \approx RzN \). All the super-nodes see the same bandwidth since all queries pass through them. If we assume that the query crosses each edge in the multicast tree (using an approach similar to [6]) then it is necessary and sufficient for the maximum degree to be 3, thus \( B_{\text{max}} = 3RzN \), which is independent of \( s \).

Since there are \( CN \) total content items, the average disk space is \( D_{\text{ave}} = CNz/N = Cz \). Since all content is stored on the supernodes, the maximum disk space is \( D_{\text{max}} = CNz/Ns = Cz/s \).

In the super-node system, each content is only copied (at most) one time. The average processing requirement is not very meaningful since like average bandwidth, no node experiences this load. Nodes either see almost no load, or maximum load. We assume a linear complexity for search, so that \( P = RD(f/z)N; P_{\text{ave}} = RD_{\text{ave}}(f/z)N = RCf/N \).

Since all the queries are processed by the super-nodes, we only need to compute the number of content items on each super-node, and then multiply by the query rate: \( P_{\text{max}} = RD_{\text{max}}(f/z)N = \frac{CN}{sN}fRN = \frac{C}{s}fN \).

4.1.2 Trade-offs and numerical values

One interesting feature of this model is that \( P_{\text{max}}B_{\text{ave}} = fzCPR^2N^2 \), which is independent of \( s \). So, there is a trade-off between average bandwidth and maximum CPU utilization. In the interest of considering some numerical values, we will set \( s = 1/\sqrt{N} \), which means each super-node has as many leaf nodes as there are super-nodes. In practice, one will probably prefer to minimize bandwidth to the extent that it is possible for the super-nodes to handle the load.

In the following table, we present performance metrics for the super-node algorithm with \( s = 1/\sqrt{N} \), \( N = 2^{19} \) and the values of \( R, C, f, z \) from Table 1.

| Symbol | Value |
|--------|-------|
| \( B_{\text{ave}} \) | \( Rz\sqrt{N} = 70B/s = 560bps \) |
| \( B_{\text{max}} \) | \( 3RzN = 150,000B/s = 1.2Mbps \) |
| \( D_{\text{ave}} \) | \( Cz = 16KB \) |
| \( D_{\text{max}} \) | \( Cz\sqrt{N} = 11MB \) |
| \( P_{\text{ave}} \) | \( RCfN = 420kFLOP/s \) |
| \( P_{\text{max}} \) | \( RCfN\sqrt{N} = 300MFLOP/s \) |

The values look reasonable. Since modern CPUs have processing power on the order of 4 GFLOPS, the above processing requirements are not more than one CPU. The figure we might be most concerned about is \( B_{\text{max}} \), however that value is independent of \( s \). Now we compare the above with the percolation search algorithm for unstructured networks.

4.2 Percolation search in power-law networks

In [9] and subsequent work, the authors show that using a combined random walk data replication/random walk query distribution scheme (to be detailed below) one can achieve sub-linear (in \( N \)) query complexity in power-law networks. Below we summarize this algorithm.

The degree distribution \( p_k \) of the network describes the probability to draw a node with degree \( k \) from the network \( p_k = Ak^{-\tau} \) where \( A \) is a normalization constant such that \( \sum_{k=2}^{k_{\text{max}}} p_k = 1 \). The main result of [9] is the following three-step algorithm: Step 1, Content
**List Implantation:** To insert cached replicas of the content, a random walk is performed and each peer visited during this random walk receives a copy of the index data. For \( \tau = 2 \), the length of the random walk should be \( O(\log N) \).

**Step 2, Query Implantation:** The above process is performed for each query. **Step 3, Bond percolation:** After the query implantation, each node that has received a query so far will forward the query to each of its neighbors with a probability \( q = \gamma q_c \), where \( q_c \) is the percolation threshold: \( \langle k \rangle / (\langle k^2 \rangle - \langle k \rangle) \), and \( \gamma \) is a small number greater than unity.

For random power-law networks, there are two parameters one may control: \( \tau \) the exponent of the power-law, and \( k_{max} \) the maximum number of neighbors any node has. In this work we will only consider \( \tau = 2 \).

For \( \tau = 2 \), \( \frac{1}{N} = \sum_{k=1}^{k_{max}} \frac{1}{k} \approx \pi^2/6 \approx 1.6 \). Next we consider the scaling of the average and maximum of bandwidth, disk and processing requirements for the percolation search.

### 4.2.1 Resource requirements

Given that a node receives a query, with probability \( q \), each neighbor sees that query. So, the total number of edges to see a query will be \( qE \), where \( E \) is the total number of edges. Since \( E = \langle k \rangle N/2 \), and \( q \approx q_c = \langle k \rangle / (\langle k^2 \rangle - \langle k \rangle) \approx \langle k \rangle / \langle k^2 \rangle \), we have that the average bandwidth cost is \( B_{ave} = \frac{RNEq}{2N} = Rz \langle k^2 \rangle N / 2k \). When \( p_k = A/k^2 \), \( \langle k \rangle = A \ln k_{max} \), \( \langle k^2 \rangle = Ak_{max} \), then we have \( B_{ave} \approx RzN/A \ln k_{max} \).

To compute the maximum bandwidth, we need to look at the highest degree node and see how many of its neighbors will see the query. The highest degree node has \( k_{max} \) neighbors, and on average \( qk_{max} \) will see the query, thus: \( B_{max} = RzNk_{max}q = RzNk_{max}q_c \approx RzN \ln k_{max} \).

For a power-law random network with exponent \( \tau = 2 \), the content is cached on \( \log_2 N \) nodes. Thus, the average storage requirements are \( D_{ave} = CzN \log_2 N = Cz \log_2 N = O(\ln N) \).

To compute the storage required for the maximum node is more involved. We model a random walk on a random network as each step selecting a random node of degree \( m \) with probability \( m p_m / \langle k \rangle \). The probability of selecting a node of the highest degree is \( P_s = \frac{k_{max}p_{k_{max}}}{q_{k_{max}}} = (k_{max} \ln k_{max})^{-1} \). We assume that we select each of the nodes of degree \( k_{max} \) with equal probability, so the probability we select each one of them is: \( Q_s = P_s (Np(k_{max}))^{-1} \). There are \( \log_2 N \) steps, so the number of content caches that make it to highest degree nodes is \( F_{max} = Q_s \log_2 N = \frac{k_{max} \ln N}{\ln 2A \ln k_{max}} \). Now we can compute the maximum storage requirements:

\[
D_{max} = CzNF_{max} = \frac{Czk_{max} \ln N}{A \ln 2 \ln k_{max}}
\]

As before, we assume that the search time is linear in the number of items stored at each node. Since we have already computed the number of items stored at each node, we have:

\[
P_{ave} = RND_{ave}(f/z) = RCfN \ln N \frac{1}{\ln 2}
\]

\[
P_{max} = RND_{max}(f/z) = RCfN \frac{k_{max} \ln N}{A \ln 2 \ln k_{max}}
\]

To reduce \( P_{max} \) we need to reduce \( k_{max} \), but that will increase \( B_{ave} \).

#### 4.2.2 Trade-offs and numerical values

In the percolation search, like the super-node system, we can decrease the average bandwidth required at the expense of increasing the maximum processor utilization. Using the percolation search \( B_{ave}P_{max} = fzCR^2 N^2 \ln k_{max}^2 / \ln N \), which is similar to the super-node architecture except with some logarithmic factors.

In order to minimize average bandwidth, we should choose \( k_{max} \) to be as large as possible. In general, the percolation search algorithm behaves like the ideal super-node algorithm with \( s \approx 1/k_{max} \).

To compare to the super-node case, we choose \( k_{max} = \sqrt{N} \), \( N = 2^{19} \) and the values of \( R, C, f, z \) from Table 1 and using the same constants we assumed in 4.1.2, which means our highest degrees are comparable to super-nodes. The results for this case are summarized in the following table:

| Parameter | Value |
|-----------|-------|
| \( B_{ave} \) | \( 4Rz\sqrt{N} \ln N \approx 70B/s = 560bps \) |
| \( B_{max} \) | \( 5RzN \ln N \approx 330KB/s = 2.7Mbps \) |
| \( D_{ave} \) | \( Cz \ln N \( \ln \ln N \) \approx 300KB \) |
| \( D_{max} \) | \( Cz\sqrt{N} \ln N \approx 52MB \) |
| \( P_{ave} \) | \( RCfN \ln N \approx 7.9MFlOp/s \) |
| \( P_{max} \) | \( RCfN \sqrt{N} \approx 1.4GFlOp/s \) |

For \( D_{max} \) and hence, \( P_{max} \) there is a constant factor overhead of \( 2/(A\ln 2) \approx 4.6 \) when compared to the ideal super-node case. For all other metrics, there is an \( O(\ln N) \) overhead for using the percolation search, however, for networks of size \( N = 2^{19} \), due to the division by a constant, the difference is not very great.

#### 4.2.3 Simulation results

Our simulations use the Netmodeler package[1]. We insert 1000 content objects at uniformly selected nodes on a power-law network with \( \tau = 2 \), and then make 1000 queries from uniformly selected nodes. We are
particularly concerned with the maximum demands made on any node. Some results are in the following table. BW is the total number of times each query is copied to search the entire network, and Max C/O is the average of the maximum CPU where 1 is the cost to evaluate the query, per content object in the network.

| Nodes | q  | ttl | Hit-rate | BW   | Max C/O |
|-------|----|-----|----------|------|---------|
| $2^{19}$ | 0.01 | 20  | 0.961    | 10,428 | 0.0075  |
| $2^{20}$ | 0.01 | 21  | 0.966    | 21,045 | 0.0042  |

We see that for the case on 1000 content objects, the maximum node had to search 7.5 and 4.2 objects for each query on average, for the cases of $2^{19}$ and $2^{20}$ nodes respectively. To scale these results up to our assumptions of 20 content objects per node. Additionally, the total query byte rate rate will be $zNR = 800 \times 2^{19} \times 1.2 \times 10^{-4} \approx 48,500 Bps = Q$.

For $N = 2^{19}$ we have:

$$B_{ave} = \frac{10,428}{2^{19}}Q = 7.7 Kbps$$

$$D_{max} = 13,280 \times 2^{19} \times \frac{7.5}{1000} = 52.21 MB$$

$$P_{max} = 20 \times 2^{19}Q \cdot \frac{332}{8001000} = 1.6 GFlOp/s$$

The above parameters are a relatively close match to the predictions of the previous section. For $N = 2^{20}$ we have $Q' = 2Q$:

$$B_{ave} = \frac{21,045}{2^{20}}Q' = 15.6 Kbps$$

$$D_{max} = 13,280 \times 2^{20} \times \frac{4.2}{1000} = 58.5 MB$$

$$P_{max} = 20 \times 2^{20}Q' \cdot \frac{332}{8001000} = 3.6 GFlOp/s$$

The other parameters such as $D_{ave}, P_{ave}$ and $B_{max}$ are not dependent on the nonlinearities of percolation, and as such match predictions of the previous section.

Our simulations verify that the average bandwidth required is much less than analog modems can provide and the maximum processing requirements are met by one modern desktop CPU.

5. Comparison of unstructured to structured image search

In order to compare unstructured search with current DHT-based approaches, we took PRISM [8] as a base for comparison. PRISM is a recent system with a clear focus on similarity queries over high-dimensional vectors.

The PRISM paper also describes load balancing between PRISM peers. However, within the following, we will consider PRISM without load balancing, as perfect load balancing would amount to all peers carrying the average load.

The performance metrics for the vanilla PRISM algorithm without load balancing. Are given in the table below. We chose visiting all 11 reference pairs for our calculation. Again $N = 2^{19}$ and the values of $R, C, f, z$ from Table 1.

$$B_{ave} = 21 \cdot \frac{RNz}{N} = 2.02 B/s = 16.1 Bps$$

$$B_{max} = \frac{1}{2} RNz = 12,600 B/s = 100 kbps$$

$$D_{ave} = 11Cz = 176 KB$$

$$D_{max} \approx 0.1 CzN \approx 840 MB$$

$$P_{ave} \approx 1.8RCfN = 670 kFlop/s$$

$$P_{max} \approx 0.25RCfN^2 = 55GFlOp/s$$

As Figs. 2 and 3, as well as Tab. 5 present our experiments with a simulation of PRISM. Without load balancing, PRISM behaves to quite an extent like a client/server system: most load hits few servers. Little
load is distributed, so there is low communication cost. For Euclidean distance, PRISM presents—if any—only small advantages over the super-peer network. In our current setup, in order to find 75% of the top 20 documents, we have to visit each vector more than one time on average, i.e. PRISM performs worse than random search. If one wants to push the recall to 100% (as good as the super peer method), each item is considered even more times on average, incurring a clear efficiency penalty with respect to a full scan.

The second finding is that the distribution of data items over peers is heavily skewed, emphasizing the need for load balancing as proposed in [8]. Our experiment used $32 \times 32 = 1024$ pairs. In these experiments, the first 5 most used pairs account for more than 10% of the traffic, the first 15 pairs account for more than 25% of the traffic, and the first 60 pairs account for more than 50% of the traffic. To highlight this fact, Tab. 5 shows PRISM without load balancing. Here, PRISM functions almost in a client/server-alike fashion. Please note that with the proper use of load balancing, PRISM would thus much behave like a super-peer network discussed in Section 4.1, but with slightly higher load for the super-peers.

The third finding, finally, should spawn a series of new experiments: The performance of systems like PRISM depends also on data set and distance measure. However, most of the distributed indexing literature is fixated on the Euclidean metric and similar distance measures. Our experiments show that it is clearly worthwhile to investigate deeper into the performance of such systems when using more diverse distance measures.

6 Conclusion

Summarizing, the performance of structured and unstructured systems seem to be pretty close in our application domain, while unstructured systems have the advantage of being more flexible with respect to the queries they allow. We should mention that this conclusion is similar to the recent paper of [16].

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