

# SCAN: A Small-World Structured P2P Overlay for Multi-Dimensional Queries

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## ABSTRACT

This paper presents a structured P2P overlay SCAN that augments CAN overlay with long links based on Kleinberg's small-world model in a  $d$ -dimensional Cartesian space. The construction of long links does not require the estimate of network size. Queries in multi-dimensional data space can achieve  $O(\log n)$  hops by equipping each node with  $O(\log n)$  long links and  $O(d)$  short links.

## Categories and Subject Descriptors

H.3.3 [Information Storage and Retrieval]: Information Search and Retrieval – Search process.

## General Terms

Algorithms, Design, Experimentation.

## Keywords

P2P, small-world, multi-dimensional queries.

## 1. INTRODUCTION

Information retrieval in large-scale distributed environments often involves multi-dimensional data management and queries. CAN overlay supports data partition and query in a  $d$ -dimensional Cartesian space [4]. It achieves  $O(dn^{1/d})$  query hops. This paper introduces SCAN that builds long links in CAN overlays based on Kleinberg's small-world model [2]. It can achieve  $O(\log n)$  hops by equipping each node with  $O(\log n)$  long links. Compared with previous small-world solution Symphony [3], SCAN approximates Kleinberg's small-world network in multi-dimensional data space without requiring estimate of network size. eCAN also achieves  $O(\log n)$  hops [6] by building express ways in CAN overlay. Long link construction depends on the joining process of nodes. In [1], small-world long links are built in Delaunay-graph-based networks. It uses a piggy-backing method in node joining process to add long links. Long links in SCAN can be built during or after the construction of the underlying CAN.

## 2. ARCHITECTURE OF SCAN

In a  $d$ -dimensional SCAN, each node is identified by a vector  $v \langle x_1, x_2, \dots, x_d \rangle$ .  $x_i$  is drawn from a real interval  $R = [0, H]$  ( $H > 1$ ). The first node holds the complete  $d$ -dimensional Space  $R^d$ . Forthcoming joining nodes split the zones of existing nodes in half along one dimension in a cyclic way. Data objects are identified by vector IDs drawn from  $R^d$  and are stored at the node the vector IDs of data objects fall in. To uniformly partition the  $d$ -

dimensional space, a joining node first draws a random ID  $v \langle x_1, x_2, \dots, x_d \rangle$ , where  $x_i$  follows the uniform distribution in  $[0, H]$ . Then, the node locates the existing node that holds this random ID and split it in one dimension. Nodes use the central point of the range they hold as their node IDs. Each node maintains  $O(d)$  short links to their neighboring nodes. Long links are added for nodes in a small-world way to speed up queries. Figure 1 shows a partial view of a 2- $d$  SCAN topology.

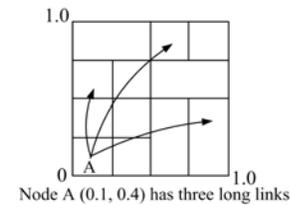


Figure 1. Topology of SCAN.

## 2.1 Building long links in SCAN

In space  $R^d$ , we define the Manhattan distance between two points  $v \langle x_1, x_2, \dots, x_d \rangle$  and  $u \langle y_1, y_2, \dots, y_d \rangle$  as  $d(v, u) = \sum_{i=1}^d d_i(x_i, y_i)$ .

$d_i(x_i, y_i) = \min\{\text{abs}(x_i - y_i), H - \text{abs}(x_i - y_i)\}$  is the coordinate distance in the  $i$ th dimension. The maximum Manhattan distance  $L_{\max}$  is  $dH/2$  because in each dimension the maximum coordinate distance is  $H/2$ .

To build long links, a node  $v \langle x_1, x_2, \dots, x_d \rangle$  first draws  $K$  real numbers  $r_1, r_2, \dots, r_K$  following the harmonic distribution in real interval  $[0, L_{\max}]$ . Then, for each  $r_i$ , a vector point  $l_i \langle y_1, y_2, \dots, y_d \rangle$  is generated as a seed ID at distance  $r_i$  from  $v$ . Finally, node  $v$  locates node  $v_i$  that is responsible for  $l_i$  and connects  $v_i$  as a long link.

Since we do not know the network size, we set  $K = \lfloor C \log_2 N \rfloor$  where  $N$  is a predefined large integer satisfying  $N \gg dn^{1/d}$  and  $C \geq 1$  is a predefined constant integer.  $r_1, r_2, \dots, r_K$  are produced by a harmonic distribution generator in  $[0, L_{\max}]$ .  $r_i = L_{\max} / 2^x$ , where  $x$  is a real number randomly drawn from the real interval  $[0, \log_2 N]$  for  $i = 1, 2, \dots, K$ .

Given a distance  $r_i$  from  $v$ , there are multiple candidate points for  $l_i$ . We randomly generate a real vector  $\tau_i \langle s_1, s_2, \dots, s_d \rangle$  so that point  $l_i \langle x_1 + s_1, x_2 + s_2, \dots, x_d + s_d \rangle$  is at distance  $r_i$  from  $v$  (plus is in wrap mode in  $[0, H]$ ). We iteratively generate  $s_k$  by regarding the remainder coordinates  $s_{k+1}, s_{k+2}, \dots, s_d$  as one coordinate. Initially, let  $M_1 = L_{\max}$ ,  $D_1 = r_i$ , and  $\delta = L_{\max} / d$ . To get  $s_k$ , following steps are repeated for  $k = 1, 2, \dots, d$ :

(STEP 1): If  $M_k \leq \delta$ , let  $s_k = M_k$  and return;

(STEP 2):  $I = [0, \delta] \cap [D_k - M_k + \delta, D_k]$ ;

(STEP 3): Get a random real number from  $I$  as  $s_k$ ;

(STEP 4):  $M_{k+1} = M_k - \delta$ ;  $D_{k+1} = D_k - s_k$ .

Long links from  $v$  can approach a remote node in two directions along one dimension. We randomly assign  $s_k$  a positive or negative sign with probability 1/2. Then  $l_i \langle y_1, y_2, \dots, y_d \rangle$  is obtained by  $y_k = x_k + s_k$  ( $k = 1, 2, \dots, d$  and plus is in wrap mode in  $[0, H]$ ). Node  $v$  locates the remote node  $v_i$  that holds  $l_i$ . After finding  $v_i$ ,  $v$  inserts it into the routing table.

Although  $K = \lfloor C \log_2 N \rfloor$  is larger than  $\log n$ , many seed IDs are actually located in the same node and the expected number of distinct long links is  $O(C \log_2 dn^{1/d})$ . It is the harmonic distribution of  $r_i$  in  $[0, L_{max}]$  that makes the long links form a small-world overlay.

## 2.2 Query routing in SCAN

The size of ranges should be considered in a greedy routing process. Let  $Z_v = \langle z_1, z_2, \dots, z_d \rangle$  be the  $d$ -dimensional range that the remote node  $v$  currently holds, where  $z_i = [zx_i, zy_i]$  is the real interval in the  $i$ th dimension that  $v$  occupies. The range Manhattan distance from node  $v$  to the target point  $t \langle y_1, y_2, \dots, y_d \rangle$  is

$$d_r(v, t) = \sum_{i=1}^d q_i(z_i, y_i) \cdot q_i(z_i, y_i)$$

in the  $i$ th dimension.  $q_i(z_i, y_i) = 0$  if  $y_i \in z_i$ . Else  $q_i(z_i, y_i) = \min\{d_i(zx_i, y_i), d_i(zy_i, y_i)\}$ . In each hop, node selects from its links the one with the shortest range Manhattan distance to the target point as the next hop. When the node at distance zero to the target point is reached, the target point is located.

When  $C = 2^d$ , the expected routing hops is bounded by  $O(\log_2 dn^{1/d})$  because each long link can help reduce the distance by half with probability  $1/2^d$ . When  $d$  is large, building  $2^d \log_2 N$  long links is prohibitive. Fortunately, when  $d > \log_2 n$ ,  $O(\log n)$  query hops can be achieved without using long links. Moreover, we can use  $K = 4 \log_2 N$  seed IDs to build long links and achieve  $O(\log_2 2n^{1/2})$  query hops in most cases as long as  $n > (d/2)^{(1/2-1/d)}$ , i.e.,  $dn^{1/d} < 2n^{1/2}$ . It is because that when  $dn^{1/d} < 2n^{1/2}$ ,  $d$ -dimensional CAN overlays of size  $n$  have shorter longest query hops than 2-d CAN overlays of the same size  $n$ . Adding the same number of long links, the  $d$ -dimensional CAN overlays can still achieve shorter query hops than the 2-d CAN overlays. Using  $K = 4 \log_2 N$  seed IDs can achieve  $O(\log_2 2n^{1/2})$  query hops in 2-d SCAN overlays. Thus, using the same number of seed IDs can also achieve  $O(\log_2 2n^{1/2})$  query hops in  $d$ -dimensional SCAN overlays of size  $n$  if  $dn^{1/d} < 2n^{1/2}$ .

## 3. EXPERIMENTS AND CONCLUSIONS

Figure 2 (a) depicts the distribution of query hops in a 2-d Kleinberg small-world mesh (Kleinberg 2D) and a 2-d SCAN with  $N = 2^{20}$  (SCAN\_2D). Both have 1024 nodes. Kleinberg's 2-d mesh is strictly regular, having shorter average query hops. Figure 2 (b) shows that in a 2-d SCAN with  $K = 4 \log_2 N$ , the average number of long links (curve 2\_avg\_rt) is bounded by  $4 \log_2(2n^{1/2})$  (curve 4LOG). The average query hops (curve 2\_avg\_qr) is bounded by  $\log_2(2n^{1/2})$  (curve LOG). Figure 2 (c) and (d) demonstrate that using  $K = 4 \log_2 N$  seed IDs in SCAN overlays with  $d = 3, 4$  and  $5$ , the average query hops (curves  $d\_avg\_qr$ ) and the maximum query hops (curves  $d\_max\_qr$ ) are bounded by those of 2-d SCAN overlays. Figure 3 shows that as  $d$  increases, CAN overlays without long links can also achieve  $O(\log n)$  query hops with routing table size of  $O(\log n)$ . Experiments demonstrate the effectiveness and the efficiency of SCAN. It can be extended

to support multi-dimensional queries based on other distance metrics. Future work also includes applying the load balancing method in one-dimensional ring [7] to SCAN.

## 4. ACKNOWLEDGEMENTS

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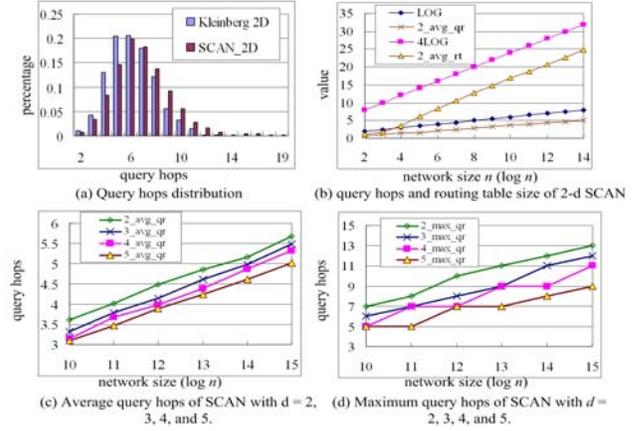


Figure 2. Topology properties of SCAN.

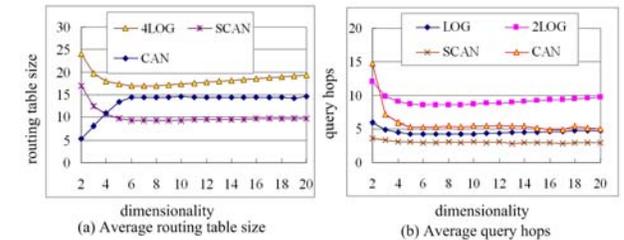


Figure 3. CAN and SCAN overlays with different dimensionality.

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