Searching for Frequent Colors in Rectangles

Marek Karpinski*  
Yakov Nekrich†

Abstract

We study a new variant of colored orthogonal range searching problem: given a query rectangle \( Q \) all colors \( c \), such that at least a fraction \( \tau \) of all points in \( Q \) are of color \( c \), must be reported. We describe several data structures for that problem that use pseudo-linear space and answer queries in poly-logarithmic time.

1 Introduction

The colored range reporting problem is a variant of the range searching problem in which every point \( p \in P \) is assigned a color \( c \in C \). The set of points \( P \) is preprocessed in the data structure so that for any given rectangle \( Q \) all distinct colors of points in \( Q \) can be reported efficiently. In this paper we consider a variant of this extensively studied problem in which only frequently occurring colors must be reported.

We say that a color \( c \in C \) \( \tau \)-dominates rectangle \( Q \) if at least a \( \tau \)-fraction of points in \( Q \) are of that color: \(|\{p \in P \cap Q \mid \text{col}(p) = c\}| \leq \tau|P \cap Q|\), where \( \text{col}(p) \) denotes the color of point \( p \). We consider several data structures that allow us to report colors that dominate \( Q \).

Motivation Standard colored range reporting problem arises in many applications. Consider a database in which every object is characterized by several numerical values (point coordinates) and some attribute (color). For instance the company database may contain information about age and salary of each employee. The attribute associated with each employee is her position. The query consists in reporting all different job types for all employees with salary between 40.000 and 60.000 who are older than 40 and younger than 60 years old. Colored range reporting also occurs naturally in computational biology applications: each amino acid is associated with certain attributes (hydrophobic, charged, etc.). We may want to report different attributes associated with amino acids in certain range [8].

However, in certain applications we are not interested in all attributes that occur in the query range. Instead, we may be interested in reporting the typical attributes. For instance, in the first example above we may wish to know all job types, such that at least a fraction \( \tau \) of all employees with a given salary and age range have a job of this type. In this paper we describe data structures that support such and similar queries.

Related Work. Traditional colored range reported queries can be efficiently answered in one, two, and three dimensions. There are data structures that use pseudo-linear space and answer one- and two-dimensional colored range reporting queries in \( O(\log n + k) \) time [6], [7] and three-dimensional colored queries in \( O(\log^2 n + k) \) time [6], where \( k \) is the number of colors. A semi-dynamic data structure of Gupta et al. [6] supports two-dimensional queries in \( O(\log^2 n + k) \) time and insertions in \( O(\log^3 n) \) amortized time. Colored orthogonal range reporting queries in \( d \) dimensions can be answered in \( O(\log n + k) \) time with a data structure that uses \( O((n^{1+\varepsilon})) \) space [1], but no efficient pseudo-linear space data structure is known for \( d > 3 \).

De Berg and Haverkort [4] consider a variant of the colored range searching in which only significant colors must be reported. A color \( c \) is significant in rectangle \( Q \) if at least a fraction \( \tau \) of points of that color belong to \( Q \): \(|\{p \in Q \cap P \mid \text{col}(p) = c\}| \geq \tau|\{p \in P \mid \text{col}(p) = c\}|\). For \( d = 1 \), de Berg and Haverkort [4] describe a linear space data structure that answers queries in \( O(\log n + k) \) time, where \( k \) is the number of significant colors. For \( d \geq 2 \) significant queries can be answered approximately: in \( O(\log n + k) \) time we can report all a set of colors such that each color is \((1 - \varepsilon)\tau\)-significant for a fixed constant \( \varepsilon \) and all \( \tau \)-significant colors are reported. The only known data structure that efficiently answers exact significance queries uses cubic space [4].

Our Results In this paper we show that we can find domination colors in an arbitrary \( d \)-dimensional rectangle in poly-logarithmic time using a pseudo-linear space data structure.

- We describe a static \( O(\tau n) \) space data structure that supports one-dimensional queries in \( O(\tau \log n \log \log n) \) time. A static \( O(\tau n \log \log n) \) space data structure supports one-dimensional domination queries in \( O(\tau \log n) \) time.
- In the case when all coordinates are integers bounded by \( U \), there is a \( O(\tau n) \) space static data structure that supports one-dimensional domination queries in \( O(\tau \log n \log \log \log U) \) time.
- There is a dynamic \( O(\tau n) \) space data structure

\*Dept. of Computer Science, University of Bonn. Email marek@cs.uni-bonn.de.
\†Dept. of Computer Science, University of Bonn. Email yasha@cs.uni-bonn.de.
\‡Further we will assume that parameter \( \tau \) is fixed and simply say that a color \( c \) dominates rectangle \( Q \).
that supports one-dimensional domination queries and insertions in \(O(\tau \log n)\) time and deletions in \(O(\tau \log n)\) amortized time. We can reduce the update time to (amortized) \(O(\log n)\) by increasing the space usage to \(O(\tau n \log n)\)

- There is a data structure that supports domination queries in \(d\) dimensions in \(O(\tau \log^d n)\) time and uses \(O(\tau n \log^{d-1} n)\) space

- There is a dynamic data structure that answers domination queries in \(d\) dimensions in \(O(\tau \log^{d+1} n)\) time, uses \(O(\tau n \log^{d-1} n)\) space, and supports insertions in \(O(\tau \log^{d+1} n)\) time and deletions in \(O(\tau \log^{d+1} n)\) amortized time.

We describe static and dynamic data structures for one-dimensional domination queries in sections 2 and 3. Data structures for multi-dimensional domination queries are described in section 4.

2 Static Domination Queries in One Dimension

The following simple property plays an important role in all data structures for domination queries.

**Observation 1** If \(Q = Q_1 \cup Q_2\), \(Q_1 \cap Q_2 = \emptyset\), and color \(c\) is dominant in \(Q\), then either \(c\) is dominant in \(Q_1\) or \(c\) is dominant in \(Q_2\).

Due to this property a query on a set \(Q\) can be decomposed into queries on some disjoint sets \(Q_1, \ldots, Q_p\) such that \(\bigcup Q_i = Q\) and \(p\) is a constant: we find the dominating colors for each \(Q_i\) and for each color \(c\) that dominates some \(Q_i\) we determine whether \(c\) dominates \(Q\) by a range counting query.

Our data structure is based on the same approach as exponential search trees [2]. Let \(P\) be the set of all points. In one-dimensional case we do not distinguish between a point and its coordinate. \(P\) is divided into \(\beta_n\) intervals \(I_1, \ldots, I_{\beta_n}\) so that each \(P_i = P \cap I_i\) contains between \(n^{2/3}/2\) and \(2n^{2/3}\) points and \(\beta_n = \Theta(n^{1/3})\). Let \(l_i\) and \(r_i\) denote the left and right bounds of interval \(I_i\). For each \(1 \leq i \leq j \leq \beta\), the list \(L_{ij}\) contains the set of colors that dominate \([l_i, r_j]\). We denote by \(n_{ij}\) the total number of points in \([l_i, r_j]\).

Each interval \(I_i\) is recursively subdivided in the same manner: an interval that contains \(m\) points is divided into \(\beta_m\) subintervals and each subinterval contains between \(m^{2/3}/2\) and \(2m^{2/3}\) points. If some interval \(I_j\) is divided into \(I_{j,1}, \ldots, I_{j,\beta_j}\), then we say that \(I_{j,1}\) is a parent of \(I_{j,1}\). The tree \(T\) reflects the division of intervals into sub-intervals: each tree node \(u\) corresponds to an interval \(I_u\) and a node \(v\) is a child of \(v\) if and only if \(I_v\) is a child of \(I_u\). The root of \(T\) corresponds to \(P\) and leaves of \(T\) correspond to points of \(P\). The height of \(T\) is \(O(\log \log n)\). For every color \(c\), we also store all points of color \(c\) in a data structure that supports range counting queries.

Consider a query \(Q = [a, b]\). Let \(l_a\) and \(l_b\) be the leaves of \(T\) in which \(a\) and \(b\) are stored, and let \(q\) be the lowest common ancestor of \(l_a\) and \(l_b\). The search procedure visits all nodes on the path from \(l_a\) to \(q\) (\(l_b\) to \(q\) for each visited node \(u\) we construct the set of colors \(S_u\), such that every \(c \in S_u\) dominates \(I_u \cap [a, b]\). We also compute the total number of points in \(I_u \cap [a, b]\). Let \(u\) be the currently visited node of \(T\) situated between \(l_b\) and \(q\), and suppose that the node \(v\) visited immediately before \(u\) is the \((i+1)\)-st child of \(u\). Due to Observation 1 only colors stored in \(L_{ij}\) may dominate \(I_u \cap Q\). For each color \(c\) in \(L_{ij}\cup S_v\) we count how many times it occurs in \(I_u \cap Q\) using the range counting data structure for that color. Thus we can construct \(S_u\) by answering at most \(2\tau\) counting queries. Nodes between \(l_a\) and \(q\) are processed in the same way. Finally, we examine all colors in \(S_v\) and \(S_a\) and list \(L_{ij}\) of the node \(q\), where \(p\) and \(r\) are nodes on the paths from \(q\) to \(l_a\) and \(l_b\) respectively, \(p\) is the \(i\)-th child of \(q\), and \(r\) is the \(j\)-th child of \(q\). The search procedure visits \(O(\log \log n)\) nodes and answers \(O(\tau \log \log n)\) counting queries. Hence, queries can be answered in \(O(\log \log n \log \log n)\) time.

If an interval \(I\) contains \(m\) points, then all lists \(L_{ij}\) contain \(O(m^{2/3})\) elements. Data structures for range counting queries use \(O(n)\) space. Therefore the space usage of our data structure is \(O(n)\).

We can reduce the query time to \(O(\log n)\) by storing range counting data structures for each interval: for every interval \(I_u\) and every color \(c\), such that \(\{ p \in P \cap I_u \mid \text{col}(p) = c \} \neq \emptyset\), we store a data structure that supports range counting queries in \(O(\log |I_u|)\). The total number of colors in all intervals \(I_u\) for all nodes \(u\) situated on the same level of tree \(T\) does not exceed the number of points in \(P\). Therefore the total number of elements in all range counting data structures is \(O(n \log \log n)\). The query is processed in the same way as described above. We must answer \(O(\tau)\) counting queries on \(I_q\), \(O(\tau)\) range counting queries on children of \(I_q\), \(O(\tau)\) range counting queries on children of children of \(I_q\), etc. Therefore the query time is \(O(\tau \log(\log |I_q|) + \log(|I_q|^{2/3}) + \log(|I_q|^{4/9}) + \ldots)) = O(\tau \sum(2/3)^i \log n) = O(\tau \log n)\).

We obtain the following result

**Theorem 1** There exists a \(O(\tau n \log \log n)\) space data structure that supports one-dimensional domination queries in \(O(\tau \log n)\) time. There exists a \(O(\tau n)\) space data structure that supports one-dimensional domination queries in \(O(\tau \log n \log \log n)\) time.

In the case when all point coordinates are integers bounded by a parameter \(U\) we can easily answer one-dimensional counting queries in \(O(\log \log U)\) time. As shown above, a domination query can be answered by answering \(O(\tau \log \log n)\) counting queries; hence, the query time is \(O(\tau \log \log n \log \log U)\). Since it is not necessary to store range counting data structures for each
interval, all range counting data structures use $O(n)$ space.

**Theorem 2** There exists a $O(\tau n)$ space data structure that supports one-dimensional domination queries in $O(\tau \log \log U \log \log n)$ time.

### 3 Dynamic Domination Queries in One Dimension

Let $T$ be a binary tree on the set of all $p \in P$. With every internal node $v$ we associate a range $rng(v) = [l_v, r_v]$, where $l_v$ is the leftmost leaf descendant of $v$ and $r_v$ is the leaf that follows the rightmost leaf descendant of $v$. $T$ is implemented as a balanced binary tree, so that insertions and deletions are supported in $O(\log n)$ time and the tree height is $O(\log n)$. In each node $v$ we store the number of its leaf descendants, and the list $L_v$; $L_v$ contains all colors that dominate $rng(v)$. For every color $c$ in $L_v$ we also maintain the number of points of color $c$ that belong to $rng(v)$. For each color $c$ there is also a data structure that stores all points of color $c$ and supports one-dimensional range counting queries.

A query $Q = [a, b]$ is answered by traversing the paths from $l_a$ to $q$ and from $l_b$ to $q$, where $l_a$ and $l_b$ are the leaves that contain $a$ and $b$ respectively, and $q$ is the lowest common ancestor of $a$ and $b$. As in the previous section, in every visited node $u$ the search procedure constructs the set of colors $S_u$, such that every $c \in S_u$ dominates $rng(v) \cap [a, b]$. Suppose that a node $v$ on the path from $l_b$ to $q$ is visited and let $u$ be the child of $v$ that is also on the path from $l_b$ to $q$. If $u$ is the left child of $v$, then $rng(v) \cap [a, b] = rng(u) \cap [a, b]$ and $S_u = S_v$. If $u$ is the right child of $v$, then $rng(v) \cap [a, b] = rng(w) \cup \{rng(u) \cap [a, b]\}$ where $w$ is a sibling of $u$. Colors that dominate $rng(w)$ are stored in $L_w$; we know colors that dominate $(rng(u) \cap [a, b])$ because $u$ was visited before $v$ and $S_u$ is already constructed. Hence, we can construct $S_v$ by examining each color $c \in L_w \cup S_u$ and answering the counting query for each color. Since one-dimensional dynamic range counting can be answered in $O(\log n)$ time, we spend $O(\tau \log n)$ time in each tree node. Nodes on the path from $l_a$ to $q$ are processed in a symmetric way. Finally we examine the colors stored in $S_{q_1}$ and $S_{q_2}$, where $q_1$ and $q_2$ are the children of $q$, and find the colors that dominate $rng(q) \cap [a, b] = [a, b]$.

When a new element is inserted/deleted, we insert a new leaf $l$ into $T$ (remove $l$ from $T$). For every ancestor $v$ of $l$, the list $L_v$ is updated.

After a new point of the color $c_p$ is inserted, the color $c_p$ may dominate $rng(v)$ and colors in $L_v$ may cease to dominate $rng(v)$. We may check whether $c_p$ must be inserted into $L_v$ and whether some colors $c \in L_v$ must be removed from $L_v$ by performing at most $\tau + 1$ range counting queries. Since a new point has $O(\log n)$ ancestors, insertions are supported in $O(\tau \log^2 n)$ time.

When a point of color $c_p$ is deleted, we may have to delete the color $c_p$ from $L_v$. We can test this by performing one counting query. However, we may also have to insert some new color $c$ into $L_v$ because the number of points stored in descendants of the node $v$ decreased by one. To implement this, we store the set of candidate colors $L'_v$; $L'_v$ contains all colors that $(\tau/2)$-dominate $rng(v)$. For each color $c \in L'_v$ we test whether $c$ became a $\tau$-dominating color after deletion. When the number of leaf descendants of the node $v$ decreased by a factor 2, we re-build the list $L'_v$. If $P_v$ is the set of leaf descendants of $v$ (that is, points that belong to $rng(v)$), then we can construct the set of distinct colors that occur in $P_v$ in $O(|P_v| \log(|P_v|))$ time. We can also find the sets of colors that $\tau$-dominate and $(\tau/2)$-dominate $rng(v)$ in $O(|P_v| \log(|P_v|))$ time. Since we re-build $L'_v$ after a sequence of at least $|P_v|/2$ deletions, re-build of some $L'_v$ incurs an amortized cost $O(\log n)$. Every deletions may affect $O(\log n)$ ancestors; hence, deletions are supported in $O(\log^2 n)$ amortized time.

We can speed-up the update operations by storing in each tree node $v$ the set of distinct colors in $P_v$, denoted by $C_v$. For each color $c \in C_v$, we store how many times points with color $c$ occur in $P_v$. When a new point $p$ is inserted/deleted, we can update $C_v$ for each ancestor $v$ of $p$ in $O(1)$ time. Using $C_v$, we can decide whether a given new color must be inserted into $L_v$ in $O(1)$ time. Using $C_v$ we can also re-build $L'_v$ in $O(|C_v|) = O(|P_v|)$ time. Hence, we can support insertions in $O(\tau \log n)$ time and deletions in $O(\log n)$ time with help of lists $C_v$. The total number of elements in all $C_v$ is $O(\tau n \log n)$.

Thus we obtain the following

**Theorem 3** There exists a $O(\tau n)$ space data structure that supports one-dimensional domination queries and insertions in $O(\tau \log^2 n)$ time and deletions in $O(\tau \log^2 n)$ amortized time. There exists a $O(\tau n \log n)$ space data structure that supports one-dimensional domination queries and insertions in $O(\tau \log n)$ time and deletions in $O(\tau \log n)$ amortized time.

### 4 Multi-Dimensional Domination Queries

We can extend our data structures to support $d$-dimensional queries for an arbitrary constant $d$ using the standard range trees [3] approach.

**Theorem 4** There exists a $O(n \log^d n)$ space data structure that supports $d$-dimensional orthogonal range domination queries in $O(\log^{d-1} n (\log \log n)^2)$ time.

We describe how we can construct a $d$-dimensional data structure if we know how to construct a $(d-1)$-dimensional data structure. A range tree $T_d$ is constructed on the set of $d$-th coordinates of all points. An arbitrary interval $[a_d, b_d]$ can be represented as a
union of $O(\log n)$ node ranges. Hence, an arbitrary $d$-dimensional query $Q = Q'_d \times [a_d, b_d]$ can be represented as a union of $O(\log n)$ queries $Q_1, \ldots, Q_t$, where $t = O(\log n)$ and $Q_i = Q_{d-1} \times \text{rng}(v_i)$ for some node $v_i$ of $T$. In each node $v$ of $T$ we store a $(d-1)$-dimensional data structure $D_v$ that contains the first $d-1$ coordinates of all points whose $d$-th coordinates belong to $\text{rng}(v)$. $D_v$ supports modified domination queries in $d-1$ dimensions: for a $(d-1)$-dimensional query rectangle $Q$, $D_v$ outputs all colors that dominate $Q \times \text{rng}(v)$. Using $D_v$, we can find (at most $\tau$) colors that dominate $Q_i = Q'_d \times \text{rng}(v_i)$. Since $Q$ is a union of $O(\log n)$ ranges $Q_i$, we can identify a set $C$ that contains $O(\tau \log n)$ candidate colors by answering $O(\log n)$ modified $(d-1)$-dimensional domination queries. As follows from Observation 1, only a color from $C$ can dominate $Q$. Hence, we can identify all colors that $\tau$-dominate $Q$ by answering $O(\tau \log n)$ $d$-dimensional range counting queries. Thus the query time for $d$-dimensional queries can be computed with the formula $q(n,d) = O(\log n)q(n,d-1) + O(\tau \log n)c(n,d-1)$, where $q(n,d)$ is the query time for $d$-dimensional domination queries and $c(n,d)$ is the query time for $d$-dimensional counting queries. We can answer $d$-dimensional range counting queries in $O(\log^{d-1} n)$ time and $O(n \log^{d-1} n)$ space [5]. We can answer one-dimensional domination queries in $O(\log n)$ time by Theorem 1. Therefore $d$-dimensional domination queries can answered in $O(\tau \log^d n)$ time.

We can apply the reduction to rank space technique [9], [5] and replace all point coordinates with labels from $[1,n]$. This will increase the query time by an additive term $O(\log n)$. Since point coordinates are bounded by $n$, we can apply Theorem 2 and answer one-dimensional domination queries in $O((\log \log n)^2)$ time using a $O(n)$ space data structure. Since the space usage grows by a $O(\log n)$ factor with each dimension, our data structure uses $O(n \log^{d-1} n)$ space.

**Theorem 5** There exists a data structure that supports domination queries in $d$ dimensions in $O(\tau \log^d n)$ time and uses $O(n \log^{d-1} n)$ space.

The same range trees approach can be also applied to the dynamic one-dimensional data structure for domination queries. Since one-dimensional dynamic domination queries can be answered in $O(\tau n \log^{d-1} n)$ time and dynamic range counting queries can be answered in $O(\log^d n)$ time and $O(n \log^{d-1} n)$ space, $d$-dimensional domination queries can be answered in $O(\log^{d+1} n)$ time, and the space usage is $O(\tau n \log^{d-1} n)$. Since updates are supported in $O(\log^2 n)$ (amortized) time in one-dimensional case and update times grow by $O(\log n)$ factor with each dimension, $d$-dimensional data structure supports updates in $O(\log^{d+1} n)$ (amortized) time.

**Theorem 6** There is a dynamic data structure that answers domination queries in $d$ dimensions in $O(\tau n \log^{d-1} n)$ time, uses $O(\tau n \log^{d-1} n)$ space, and supports insertions in $O(\tau \log^d n)$ time and deletions in $O(\tau \log^d n)$ amortized time.

**Conclusion**

We presented data structures for a new variant of colored range reporting problem. Our data structures use pseudo-linear space and report all $\tau$-dominating colors in poly-logarithmic time in the case when the parameter $\tau$ is small, i.e. constant or poly-logarithmic in $n$. It would be interesting to construct efficient data structures for larger values of $\tau$.

**Acknowledgment**

We would like to thank Mark de Berg and Herman Haverkort for stimulating discussions and for suggestions concerning the new variant of colored range searching problem.

**References**

[1] P. K. Agarwal, S. Govindarajan, S. Muthukrishnan, Range Searching in Categorical Data: Colored Range Searching on Grid, Proc. ESA 2002, pp. 17-28.

[2] A. Andersson, M. Thorup, Dynamic Ordered Sets with Exponential Search Trees, J. ACM 54(3):13(2007).

[3] J. L. Bentley, Multidimensional Divide-and-Conquer, Commun. ACM 23(1980), 214-229.

[4] M. de Berg, H. J. Haverkort, Significant-Presence Range Queries in Categorical Data, Proc. WADS 2003, pp. 462-473.

[5] B. Chazelle, A Functional Approach to Data Structures and its Use in Multidimensional Searching, SIAM J. on Computing 17(1988), 427-462.

[6] D. Gupta, R. Janardan and M. Smid, Further Results on Generalized Intersection Problems: Counting, Reporting, and Dynamization, J. of Algorithms 19(1995), 282-317.

[7] R. Janardan, M. Lopez, Generalized Intersection Searching Problems, Internat. J. Comput. Geom. Appl. 3 (1993), 39-69.

[8] N. Madhusudhanan, P. Gupta, A. Mitra, Efficient Algorithms for Range Queries in Protein Sequence Analysis, Proc. CCCG 2005, pp. 146-149.

[9] M. H. Overmars, Efficient Data Structures for Range Searching on a Grid, J. Algorithms 9(1988), 254-275.