Algorithm for exact evaluation of bivariate two-sample Kolmogorov-Smirnov statistics in $O(n \log n)$ time.

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Abstract

We propose an $O(n \log n)$ algorithm for evaluation of bivariate Kolmogorov-Smirnov statistics for $n$ two dimensional data points. It offers few orders of magnitude of speedup over existing implementations for $n > 10^5$ samples of input. Algorithm is based on static Binary Search Trees and sweep algorithm. We share tested C++ implementation with Python bindings.

1 Introduction

1.1 Background

Kolmogorov test introduced by [1] is notable statistical procedure used for testing for difference between univariate probability distributions.

A generalization of Kolmogorov statistics to bivariate distributions was proposed by [2] with $O(n^2)$ algorithm for evaluating it for empirical distributions with $N$ samples. Procedure found numerous application in empirical research as many references to [2] indicate. C implementation of algorithm itself can be found in popular computing textbook [3]. Square time complexity however limits application of the procedure to small numbers of samples. We present $O(n \log n)$ algorithm able to process millions of samples within few minute on office PC. We hope that presented methodology will allow to better leverage the procedure for research work as well as modern big data applications.

1.2 Bivariate Kolmogorov statistics

Consider random variables $X_1$, $Y_1$, $X_2$ and $Y_2$. In accordance to [2] two-dimensional two-sample Kolmogorov-Smirnov test statistics is defined as follows:

$$D = \sup_{x,y} \| \Pr(X_1 < x, Y_1 < y) - \Pr(X_2 < x, Y_2 < y) \|$$

(1)

One can evaluate $D$ for empirical distribution function of bivariate data samples $A_1 = \{(x_1, y_1), \ldots, (x_n, y_n)\}$, $A_2 = \{(x'_1, y'_1), \ldots, (x'_n, y'_n)\}$:

$$F_i(u, v) = \frac{1}{\|A_i\|} \| \{(x, y) \in A_i : x < u, y < v \} \|$$

(2)
\[
D_{12} = \sup_{x,y} \|F_1(x, y) - F_2(x, y)\| \quad (3)
\]

\(F_1(c, y)\) for any \(c\) is constant everywhere except at \(y \in \{y_1, \ldots, y_n\}\), similar argument holds for \(F_1(x, c)\). Based on this key observation \([\quad]\) shows an \(O((n + n')^2)\) algorithm for finding exact global maximum of (3). We reproduce it here as it is important to our idea:

**Lemma 1.1.** To find \(D_{12}\) for two bivariate real samples \(A_1 = \{(x_1, y_1), \ldots, (x_n, y_n)\}, A_2 = \{(x'_1, y'_1), \ldots, (x'_n, y'_n)\}\) it is sufficient to evaluate \(\|F_1(x, y) - F_2(x, y)\|\) for all \(c \in C\), \(C = C_1 \times C_2\), \(C_1 = \{x_1, \ldots, x_n, x'_1, \ldots, x'_n\}\), \(C_2 = \{y_1, \ldots, y_n, y'_1, \ldots, y'_n\}\).

**Proof.** Let \(D(x, y) = \|F_1(x, y) - F_2(x, y)\|\) Let \((u, v) = \arg \max_{x,y} D(x, y)\), case of \((u, v) \in C\) is trivial, so we assume otherwise. let \(x_{up} = \arg \max_x \{x \in C_1 : x < u\}\) and \(y_{up} = \arg \max_y \{y \in C_2 : y < v\}\). By the definition \((x_{up}, y_{up}) \in C\) and by the fact that \(C\) is a product set we have:

\[
\exists_{(x, y) \in C}(x_{up} < x < u) \lor (y_{up} < y < u).
\]

Since \(A_1 \subset C\) and \(A_2 \subset C\) (2) implies that \(D(u, v) = D(x_{up}, y_{up})\). Thus \(\sup_{(x,y) \in C} D(x, y) = D_{12}\) \(\square\)

## 2 Evaluating extrema of expanding sum with BST.

### 2.1 Binary search trees.

A binary search tree (BST) \([4]\) is a data structure comprised of nodes satisfying following:

- Each node contains a single key and references to up to two other nodes. We call them parent node and child nodes respectively.

- A left (right) child is a child node with key smaller (greater) than parent’s key. Each node can have at most one left child and at most one right child.

- For non empty BST there is exactly one node with no parent, we call it root.

A path is a series of nodes \(n_1, \ldots, n_k\) s. t. for any \(i\) \(n_i\) is a parent of \(n_{i+1}\) and \(n_k\) is a leaf i.e. a node without children. A height of a tree \(h\) is the length of longest path in a tree. If for nodes \(n_i, n_j\) exist a path that contains both of them, and \(n_i\) precedes \(n_j\), we call \(n_i\) ancestor to \(n_j\) and \(n_j\) a descendant to \(n_i\). Each ancestor with all it’s descendants forms another BST we call subtree.

Lookup, insertion and deletion in BST can be done in \(O(h)\) operations by straightforward top-down transversal. One can put any \(N\) elements in a tree with \(h \leq 1 + \log N\), and algorithms exists to perform insertion or deletion in \(O(h)\) time while making sure that we keep \(h \leq 2\log N\). Those are of particular interest in practical application.
2.2 Keeping track of expanding sum extrema

Consider keys $k_i \in K \subset \mathbb{R}$, and values $v_i$, $i \in \{1, 2, ..., n\}$ associated with keys. For a finite ordered set $A$ with $m = \inf A$ are interested in following quantities:

$$U_A = \sup_{i \in A} \sum_{j=m}^{i} v_j, D_A = \inf_{i \in A} \sum_{j=m}^{i} v_j$$

(4)

Our goal here to be able to change values $v_i$ while keeping track of $U, D$.

Consider balanced BST s.t. node $n_i$ has key $k_i$ and keeps value $v_i$. Let $S_i$ be set of indices of descendants of node $n_i$ and $S'_i = S_i \cup \{n_i\}$. Let $r(i)$ be right child index (if exists) and $l(i)$ left child $i$ (if exists). Additionally each node keeps track of $s_i = v_i + \sum_{j \in S_i} v_j$ - sum of values in a node and its descendents.

Additionally we define $u_i, d_i$ as follows:

- if $n_i$ is a leaf:
  $$u_i = d_i = v_i$$

  (5)

- otherwise if $l(i)$ and $r(i)$ exist:
  $$u_i = \max\{s_{l(i)} + v_i, u_{l(i)}, s_{l(i)} + v_i + u_{r(i)}\};$$
  $$d_i = \min\{s_{l(i)} + v_i, d_{l(i)}, s_{l(i)} + v_i + d_{r(i)}\};$$

  (6)

- otherwise if only $l(i)$ exists:
  $$u_i = \max\{s_{l(i)} + v_i, u_{l(i)}\}, d_i = \min\{s_{l(i)} + v_i, d_{l(i)}\};$$

  (7)

- otherwise if only $r(i)$ exists:
  $$u_i = \max\{v_i, v_i + u_{r(i)}\}, d_i = \min\{v_i, v_i + d_{r(i)}\}. $$

  (8)

**Lemma 2.1.** For each $i$, $U_{S'_i(i)} = u_i$ and $D_{S'_i(i)} = d_i$, i.e. $u_i$ is maximum and $d_i$ is minimum of expanding sum of values in a subtree $S'_i(i)$.

**Proof.** For $n_i$ being single leaf this follows straight from the definition (5). Now consider a node $n_i$ s.t. it has left and right child and we know: maxima and minima of expanding sum, and total sum of values in left and right child. To find $U_{S'_i(i)}$ we consider three cases:

- Maximum occurs at $j > i$, i.e. in right subtree. All values of $n_i$ expanding sum in the right subtree are sum of $s_{l(i)}$, $v_i$ and respective values of $n_{r(i)}$ expanding sum.

- Maximum occurs at $j = i$. Expanding sum value at $j$ is $s_{l(i)} + v_i$.

- Maximum occurs at $j < i$, then $U_i(S'_i(i)) = U_i(S'_i(l(i)))$, and we assume to know r.h.s.

By evaluating these cases and picking greatest value we reproduce formulae (6), (7) and (8). Identical argument applies to minimum $D_{S'_i(i)}$. Application of induction in the upwards direction (from leaves to consecutive ancestors) finishes the proof. \qed
Using Lemma 2.1 allows us to define a data structure we will call **expanding sum extrema tree (ESET)**, that for a given series of pairs of keys \(k_i\) and values \(v_i\) keeps track of maximum and minimum of cumulative sum of values (keys are used for labelling and ordering). It is based on static binary search tree ordered by \(k_i\) and with values \(v_i\) added to nodes. If a key is repeated then two or more pairs share a node, and its value is a sum of all key values. Moreover it is supposed to have following properties:

- It can be constructed for \(N\) pairs in \(O(N \log N)\) time.
- Any key-value pair can be removed in \(O(\log N)\) time.

First property can be done by construction of balanced static binary search tree and then evaluation of \(u_i\) and \(d_i\) in bottom up manner. Second is allowed by Lemma 2.1 - as we remove value we just subtract it from node value and tree structure is unchanged. Then we need only to update \(O(\log N)\) ancestors of a given node. Obviously we could also insert values for any key that is present in a tree, but it is not any more practical than removal. General insertion with unchanged asymptotic complexity would require some additional mechanism such as AVL rotations and the structure described in Lemma 2.1 would need to become invariant or easily reproducible under such rotations. This seems to be not very relevant in practical applications and simultaneously a difficult problem beyond the scope of this work.

### 2.3 Algorithm for 2D Kolmogorov statistics

We reformulate our initial problem as follows: We have a set of triples:

\[
S = \{(x_i, y_i, v_i) \text{ for } i \in 1...N\}
\]

and we are interested in finding \(D\) such that:

\[
D = \sup_{(a', b')} \left\| \sum_{i=0}^{N} 1_{x_i < a' \land y_i < b'} v_i \right\| \tag{9}
\]

One can reproduce problem of finding 2d Kolmogorov statistics between two data samples \(T = \{(X_i, Y_i) \text{ for } i \in 1...N\}, T' = \{(X'_i, Y'_i) \text{ for } i \in 1...N'\}\) by defining following set of triples:

\[
S' = \{(X_i, Y_i, \frac{1}{N_1}) : i \in 1...N_1\} \cup \{(X'_i, Y'_i, -\frac{1}{N_2}) : i \in 1...N_2\}, \tag{10}
\]

then \(D(S')\) is exactly 2D Kolmogorov statistics for \(T_1, T_2\). We propose an Algorithm 1 to evaluate \(D\) in \(O(N \log N)\) time for \(N = |S|\). The algorithm resembles a standard procedure of computational geometry known as sweep [4] - we
Data: \( S' = \{(x_i, y_i, v_i) : i \in 1...N\} \)

Result: \( D \)

\( A \leftarrow \) ESET tree built of \((x_i, v_i)\);

\( \hat{x}_i, \hat{y}_i, \hat{v}_i \leftarrow x_i, y_i, v_i \) ordered by \( y_i \);

\( D \leftarrow 0 \);

\( j \leftarrow N \);

while \( j \geq 1 \) do

\( D = \max(D, A.\text{get\_max}) \);

\( A.\text{remove}(\hat{x}_j, \hat{v}_j) \);

\( j \leftarrow j - 1 \);

while \( j > 1 \land \hat{y}_j = \hat{y}_{j-1} \) do

\( A.\text{remove}(\hat{x}_j, \hat{v}_j) \);

\( j \leftarrow j - 1 \);

end

end

Algorithm 1: Evaluation of (9) formula.

Lemma 2.2. For a given main loop step with fixed \( j = j_0, j_0 \in \{1...N\} \)

\[ A.\text{get\_max}() = \sup_{(x,y) \in \{\hat{y}_{j_0}\} \times \{\hat{x}_1...\hat{x}_N\}} D(x,y) \]

Proof. In each step we remove one sample or more (in case of equal \( y \) values) from ESET tree in \( y \)-descending order (by \( \hat{x}_i, \hat{y}_i \) definition). Thus at step \( j_0 \) ESET-tree \( A \) contains up to date maximum of cumulative of \( v_i \) in \( x \)-ascending order for \( i \) s.t. \( \hat{y}_i < \hat{y}_{j_0} \) and this is exactly (11).

Theorem 2.3. Algorithm 1 correctly evaluates (9).

Proof. Let \( C = \{x_1...x_N\} \times \{y_1...y_N\} \). It follows from Lemma 1.1 that it suffices to evaluate \( \| \sum_{i=0}^{N} I_{x_i < a' \land y_i < y' v_i} \| \) for any \( c \in C \) to find \( D \), and it follows from \( C = \bigcup_{i} \{\hat{y}_i\} \times \{\hat{x}_1...\hat{x}_N\} \) and Lemma 2.2 that algorithm does this evaluation.

Algorithm takes \( O(N \log N) \) time and \( O(N) \) memory. Initially we need to build ESET-tree and store it in memory which takes \( O(N \log N) \) time and \( O(N) \) memory. Then we sort and store \( x_i, y_i \) and \( v_i \) by \( y_i \) which needs linear additional memory and \( O(N \log N) \) time if we use e.g. mergesort. Final loop comprises update to \( D \) which is \( O(1) \) and \( N \) removals that cost \( O(\log N) \) each and using \( O(\log N) \) memory.

3 Implementation

We share open source C++ implementation of the algorithm with bindings for Python/Numpy. Implementation was inter allia tested for agreement with \( O(n^2) \) algorithm from [3] on few sets of random real numbers, ranging from 100 to 10000. Implementation’s performance profiling confirmed significant improvement in asymptotic complexity and linear random-access-memory usage.
References


